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(54) Title: HUMAN STROKE GENE

(57) Abstract: A role of the human PDE4D gene in stroke is disclosed. Methods for diagnosis, prediction of clinical course and treatment for stroke using polymorphisms in the PDE4D gene are also disclosed.

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HUMAN STROKE GENE

RELATED APPLICATION

This is a continuation of U.S. Application _____ (2345.2010-003), which was filed on February 4, 2002, which is a continuation-in-part of U.S.

5   Application No. 09/811,352, filed March 19, 2001. The entire teachings of the above applications are incorporated herein by reference.

BACKGROUND OF THE INVENTION

Stroke is a major health problem in western societies. It is the leading cause of disability, the second leading cause of dementia and the third most common cause

10   of death (Bonita, R., *Lancet 339*:342 (1992)). As it is more common in the elderly, the public health impact of stroke will increase in the next decades with growing life expectancy. Almost 1 out of 4 men and nearly 1 out of 5 women aged 45 years will have a stroke if they live to their 85th year (Bonita, R., *Lancet 339*:342 (1992)). Strategies to diminish the impact of stroke includes prevention and treatment with

15   thrombolytics and possibly neuroprotective agents. The success of preventive measures will depend on the identification of risk factors and means to modulate their risk.

The clinical phenotype of stroke is complex but can be broadly divided into ischemic and hemorrhagic stroke. The majority of strokes (80 to 90%) are ischemic,

20   caused by obstruction of blood flow through extra- or intracranial vessels (Mohr, J.P., *et al.*, *Neurology*, *28*:754-762 (1978); Caplan, L.R., *In Stroke, A Clinical Approach* (Butterworth-Heinemann, Stoneham, MA, ed 3, 1993)). The remainder are hemorrhagic strokes (10-20%), resulting from ruptures of intracranial vessels. Ischemic stroke can be further subdivided into large vessel occlusive disease, small

25   vessel occlusive disease, and cardiogenic stroke. Transient ischemic attack (TIA), although not defined as a stroke because the signs and symptoms (which are the same as for stroke) last for a short period of time (less than 24 hours, usually 5 to 20

minutes), indicates a serious underlying risk that a stroke may follow, and it is believed that the same pathophysiologic mechanisms are responsible for TIA and ischemic stroke (Caplan, L.R., *In Stroke, A Clinical Approach* (Butterworth-Heinemann, Stoneham, MA, ed 3, 1993)).

5   The predominant risk factor for all types of stroke is hypertension (Thompson, D.W. and A.J. Furlan, *Neurosurg. Clin. N. Am.,* *8*:265-269 (1997); Agnarsson, U., *et al.,* *Ann. Intern. Med.,* *130*:987 (1999)) . Hypertension is in itself a complex disease as are the other known secondary risk factors, diabetes and hyperlipidemia. In addition, there are environmental risk factors such as smoking.

10   Stroke is therefore considered to be a highly complex disease consisting of a group of heterogeneous disorders with multiple risk factors, genetic and environmental.

The identification of genetic determinants of common diseases such as stroke, which may result from an interplay among multiple genes and between genes and environmental risk factors, has proven to be a difficult task. Studies of the

15   genetic contribution to stroke have mainly focused on rare Mendelian diseases where stroke is a part of the phenotype or on finding association with possible candidate genes such as genes contributing to hypertension or lipid metabolism. Several genes have been identified that play roles in the pathogenesis of rare stroke syndromes such as the *Notch3* gene in CADASIL (cerebral autosomal dominant arteriopathy

20   with subcortical infarctions and leukoencephalopathy) (Tournier-Lasserve, E., *et al.,* *Nat. Genet.,* *3*:256-259 (1993); Joutel, A., *et al., Nature, 383*:707 (1996)), *Cystatin C* in the Icelandic type of hereditary cerebral hemorrhage with amyloidosis (Palsdottir, A., *et al., Lancet, 2*:603-604 (1998)), *APP* in the Dutch type of hereditary cerebral hemorrhage (Levy, E., *et al., Science, 248*:1124 (1990)), and the

25   *KRIT1* gene in patients with hereditary cavernous angioma (Gunel, M., *et al., Proc. Natl. Acad. Sci. U.S.A., 92*:6620-6624 (1995); Laberge-le Couteulx, S., *et al., Nat. Genet. 23*:189 (1999); Sahoo, T., *et al., Hum. Mol. Genet. 8*:2325 (1999)) .

In addition to family history information for stroke, it is desirable to develop diagnostic methods for the early diagnosis of the disease or predisposition for the

30   development of stroke. Better means for predicting and identifying stroke should lead to better prophylactic and treatment regimens.

SUMMARY OF THE INVENTION

As described herein, it has been discovered that the gene that encodes phosphodiesterase 4D (hereinafter referred to as "PDE4D") has been correlated through human linkage studies to stroke, particularly ischemic strokes and transient

5 ischemic attacks. Five new exons, here referred to as 4D7-1, 4D7-2, 4D7-3, 4D6 and 4D8 have been identified. Three novel splice variants have also been identified (see Fig. 4).

The present invention relates to isolated nucleic acid molecules comprising the PDE4D gene. In one embodiment, the isolated nucleic acid molecule comprises

10 a nucleotide sequence selected from the group consisting of SEQ ID NO: 1 which may optionally comprise at least one polymorphism as shown in Tables 9 and 10, and the complement thereof. The invention further relates to a nucleic acid molecule which hybridizes under high stringency conditions to a nucleotide sequence selected from the group consisting of SEQ ID NO: 1 which may optionally

15 comprise at least one polymorphism as shown in Tables 9 and 10, and the complement thereof. The invention additionally relates to isolated nucleic acid molecules (e.g., cDNA molecules) encoding a PDE4D polypeptide (e.g., encoding SEQ ID NO: 2, 3, 4, 5, 6, 7, 8, 9, 10, 12 or 14 or another splicing variant of PDE4D polypeptide which includes a polymorphic site and/or novel exon selected from the

20 group consisting of 4D6, 4D7-1, 4D7-2, 4D7-3 and 4D8).

The invention further provides a method for assaying a sample for the presence of a nucleic acid molecule comprising all or a portion of PDE4D in a sample, comprising contacting said sample with a second nucleic acid molecule comprising a nucleotide sequence encoding a PDE4D polypeptide (e.g., SEQ ID

25 NO: 1 or the complement of SEQ ID NO: 1 which may optionally comprise at least one polymorphism as shown in Tables 9 and 10; a nucleotide sequence encoding SEQ ID NO: 2, 3, 4, 5, 6, 7, 8, 9, 10, 12 or 14 which may optionally comprise at least one polymorphism as shown in Tables 9 and 10, or another splicing variant of PDE4D polypeptide which includes a polymorphic site and/or exon selected from

30 the group consisting of 4D6, 4D7-1, 4D7-2, 4D7-3 and 4D8), or a fragment or derivative thereof, under conditions appropriate for selective hybridization. The

invention additionally provides a method for assaying a sample for the level of expression of a PDE4D polypeptide, or fragment or derivative thereof, comprising detecting (directly or indirectly) the level of expression of the PDE4D polypeptide, fragment or derivative thereof.

5   The invention also relates to a vector comprising an isolated nucleic acid molecule of the invention operatively linked to a regulatory sequence, as well as to a recombinant host cell comprising the vector. The invention also provides a method for preparing a polypeptide encoded by an isolated nucleic acid molecule described herein (an PDE4D polypeptide), comprising culturing a recombinant host cell of the
10   invention under conditions suitable for expression of said nucleic acid molecule.

The invention further provides an isolated polypeptide encoded by isolated nucleic acid molecules of the invention (e.g., PDE4D polypeptide), as well as fragments or derivatives thereof. In a particular embodiment, the polypeptide comprises the amino acid sequence of SEQ ID NO: 2, SEQ ID NO: 3, SEQ ID NO:
15   4, SEQ ID NO: 5, SEQ ID NO: 6 , SEQ ID NO: 7, SEQ ID NO: 8, SEQ ID NO: 9, SEQ ID NO: 10, SEQ ID NO: 12 or SEQ ID NO. 14 and containing at least one polymorphism described herein, particularly a polymorphism in all or a portion of exon 4D1, such as a SNP at 1,591,306, or one or a combination of SNPs in Table 5B. In another embodiment, the polypeptide is another splicing variant of an
20   PDE4D polypeptide, particularly a splicing variant containing all or a portion of exon selected from the group consisting of, 4D7-1, 4D7-2, 4D7-3 and 4D8. The invention also relates to an isolated polypeptide comprising an amino acid sequence which is greater than about 90 percent identical to the amino acid sequence of SEQ ID NO: 2, SEQ ID NO: 3, SEQ ID NO: 4, SEQ ID NO: 5, SEQ ID NO: 6, SEQ ID
25   NO: 7, SEQ ID NO: 8, SEQ ID NO: 9, SEQ ID NO:10, SEQ ID NO: 12 or SEQ ID NO: 14 and containing at least one polymorphism described herein, particularly a polymorphism in all or a portion of exon 4D1, such as a SNP at 1,591,306, or one or a combination of SNPs in Table 5B; preferably about 95 percent identical.

The invention also relates to an antibody, or an antigen-binding fragment
30   thereof, which selectively binds to a polypeptide of the invention, as well as to a method for assaying the presence of a polypeptide encoded by an isolated nucleic

acid molecule of the invention in a sample, comprising contacting said sample with an antibody which specifically binds to the encoded polypeptide.

The invention further relates to methods of diagnosing a predisposition to stroke. The methods of diagnosing a predisposition to stroke in an individual

5   include detecting the presence of a mutation in PDE4D, as well as detecting alterations in expression of an PDE4D polypeptide, such as the presence of different splicing variants of PDE4D polypeptides. The alterations in expression can be quantitative, qualitative, or both quantitative and qualitative. The methods of the invention allow the accurate diagnosis of stroke at or before disease onset, thus

10   reducing or minimizing the debilitating effects of stroke.

The invention additionally relates to an assay for identifying agents which alter (e.g., enhance or inhibit) the activity or expression of one or more PDE4D polypeptides. For example, a cell, cellular fraction, or solution containing an PDE4D polypeptide or a fragment or derivative thereof, can be contacted with an

15   agent to be tested, and the level of PDE4D polypeptide expression or activity can be assessed. The activity or expression of more than one PDE4D polypeptides can be assessed concurrently (e.g., the cell, cellular fraction, or solution can contain more than one type of PDE4D polypeptide, such as different splicing variants, and the levels of the different polypeptides or splicing variants can be assessed).

20   In another embodiment, the invention relates to assays to identify polypeptides which interact with one or more PDE4D polypeptides. In a yeast two-hybrid system, for example, a first vector is used which includes a nucleic acid encoding a DNA binding domain and also an PDE4D polypeptide, splicing variant, or fragment or derivative thereof, and a second vector is used which includes a

25   nucleic acid encoding a transcription activation domain and also a nucleic acid encoding a polypeptide which potentially may interact with the PDE4D polypeptide, splicing variant, or fragment or derivative thereof (e.g., a PDE4D polypeptide binding agent or receptor). Incubation of yeast containing both the first vector and the second vector under appropriate conditions allows identification of polypeptides

30   which interact with the PDE4D polypeptide or fragment or derivative thereof, and thus can be agents which alter the activity of expression of an PDE4D polypeptide.

Agents that enhance or inhibit PDE4D polypeptide expression or activity are also included in the current invention, as are methods of altering (enhancing or inhibiting) PDE4D polypeptide expression or activity by contacting a cell containing PDE4D and/or polypeptide, or by contacting the PDE4D polypeptide, with an agent

5   that enhances or inhibits expression or activity of PDE4D or polypeptide.

Additionally, the invention pertains to pharmaceutical compositions comprising the nucleic acids of the invention, the polypeptides of the invention, and/or the agents that alter activity of PDE4D polypeptide. The invention further pertains to methods of treating stroke, by administering PDE4D therapeutic agents,

10   such as nucleic acids of the invention, polypeptides of the invention, the agents that alter activity of PDE4D polypeptide, or compositions comprising the nucleic acids, polypeptides, and/or the agents that alter activity of PDE4D polypeptide.

BRIEF DESCRIPTION OF THE DRAWINGS

The foregoing and other objects, features and advantages of the invention

15   will be apparent from the following more particular description of preferred embodiments of the invention, as illustrated in the accompanying drawings.

Figs. 1A and 1B show two family pedigrees each affected by several of the stroke subtypes, including hemorrhagic stroke.

Figs. 2A, 2B and 2C show the genetic, combined and physical maps for

20   locating the PDE4D gene using 30 polymorphic markers. For the combined map, all markers have been assigned in the genetic and physical map unless otherwise indicated. (* indicates markers only assigned in physical map; ** indicates markers only assigned in genetic map).

Fig. 3 shows the genetic map of the stroke locus with exons and polymorphic

25   markers indicated. Markers identified by asterisks show association. The area defined by one drop in lod is approximately 4.6 Mb (approximately 5-6 cM).

Fig. 4 shows schematic representations of PDE4D splice variants. Splice variants 4D6, 4D7 and 4D8 are novel, as well as exons 4D6, 4D7-1, 4D7-2, 4D7-3 and 4D8. Splice variants 4DN1, 4DN2 and 4DN3 (Miro, *et al., Biochem. Biophys.*

Res. Comm., 274:415-421 (2000)), and 4D1, 4D2, 4D3, 4D4 and 4D5 (Bolger *et al.*,

Biochem. J., Pt2:539-548 (1997) are known.

      Fig. 5 is a schematic representation of the genetic map showing

microsatellites and SNP haplotypes within the stroke gene.

5   Figs. 6.1 to 6.351 show the genomic sequence of the human PDE4D gene.

      Figs. 7.1 to 7:10 show the amino acid sequences for the isoforms of the

PDE4D gene. SEQ ID NO: 2 is D4; SEQ ID NO: 3 is N2; SEQ ID NO: 4 is D5;

SEQ ID NO: 5 is N3; SEQ ID NO: 6 is D3; SEQ ID NO: 7 is N1; SEQ ID NO: 8 is

D6; SEQ ID NO: 9 is D1; and SEQ ID NO: 10 is D2.

10   Figs. 8A and 8B list all publically available PDE4D2 mRNA's and novel

eDNA segments identified by deCODE genetics.

DETAILED DESCRIPTION OF THE INVENTION

      Extensive genealogical information for a population with population-based

lists of patients has been combined with powerful genome sharing methods to map

15   the first major locus in common stroke. A genome wide scan on patients, related

within 6 meiotic events, diagnosed with stroke (ischemic and TIA) and their

unaffected relatives has been completed. Locus *STRK1* on chromosome 5q12 has

been identified through linkage studies to be associated with stroke. This locus does

not correspond to known susceptibility loci for stroke or its risk factors (such as

20   diabetes, hyperlipidemia and hypertension), and represents the first mapping of a

gene for common stroke. Until now there have been no known linkage studies of

stroke in humans showing any connection to this region of the chromosome. Based

on the linkage studies conducted, Applicants have discovered a direct relationship

between the PDE4D gene and stroke. Although the PDE4D gene (i.e., cDNA but

25   not the genomic sequence) from normal individuals is known, there have been no

studies directly investigating PDE4D and stroke. Moreover, there have been no

variant forms reported that have been associated with stroke. The full sequence of

the PDE4D gene and splice variants are reported herein. Additional single

nucleotide polymorphisms are reported in Tables 9 and 10 and may not be shown in

30   SEQ ID NO: 1.

NUCLEIC ACIDS OF THE INVENTION

       Accordingly, the invention pertains to an isolated nucleic acid molecule comprising the human PDE4D gene having at least one nucleotide alteration and correlated with incidence of stroke. The term, "PDE4D or variant PDE4D", as used

5   herein, refers to an isolated nucleic acid molecule on chromosome 5q12 having at least one altered nucleotide that is associated with a susceptibility to a number of stroke phenotypes, and also to a portion or fragment of the isolated nucleic acid molecule (e.g., cDNA or the gene) that encodes PDE4D polypeptide (e.g., the polypeptide having SEQ ID NO: 2, 3, 4, 5, 6, 7, 8, 9, 10, 12 or 14, optionally

10   comprising at least one SNP as set forth in Tables 9 and 10, or another splicing variant of a PDE4D polypeptide). In a preferred embodiment, the isolated nucleic acid molecule comprises SEQ ID NO:1 (shown in Appendix I) or the complement thereof. In another embodiment, the isolated nucleic acid molecule comprises the sequence of SEQ ID NO: 1 or the complement of SEQ ID NO: 1, except that one or

15   more single nucleotide polymorphisms as shown in Tables 9 and 10 are also present. In another embodiment, the isolated nucleic acid molecules comprises exon 4D6, 4D7-1, 4D7-2, 4D7-3 and 4D8.

       The isolated nucleic acid molecules of the present invention can be RNA, for example, mRNA, or DNA, such as cDNA and genomic DNA. DNA molecules can

20   be double-stranded or single-stranded; single stranded RNA or DNA can be either the coding, or sense, strand or the non-coding, or antisense, strand. The nucleic acid molecule can include all or a portion of the coding sequence of the gene and can further comprise additional non-coding sequences such as introns and non-coding 3' and 5' sequences (including regulatory sequences, for example). Additionally, the

25   nucleic acid molecule can be fused to a marker sequence, for example, a sequence that encodes a polypeptide to assist in isolation or purification of the polypeptide. Such sequences include, but are not limited to, those which encode a glutathione-S-transferase (GST) fusion protein and those which encode a hemagglutinin A (HA) polypeptide marker from influenza.

30   An "isolated" nucleic acid molecule, as used herein, is one that is separated from nucleic acids which normally flank the gene or nucleotide sequence (as in

genomic sequences) and/or has been completely or partially purified from other

transcribed sequences (e.g., as in an RNA library). For example, an isolated nucleic

acid of the invention may be substantially isolated with respect to the complex

cellular milieu in which it naturally occurs, or culture medium when produced by

5 recombinant techniques, or chemical precursors or other chemicals when chemically

synthesized. In some instances, the isolated material will form part of a composition

(for example, a crude extract containing other substances), buffer system or reagent

mix. In other circumstances, the material may be purified to essential homogeneity,

for example as determined by PAGE or column chromatography such as HPLC.

10 Preferably, an isolated nucleic acid molecule comprises at least about 50, 80 or 90%

(on a molar basis) of all macromolecular species present. With regard to genomic

DNA, the term "isolated" also can refer to nucleic acid molecules which are

separated from the chromosome with which the genomic DNA is naturally

associated. For example, the isolated nucleic acid molecule can contain less than

15 about 5 kb, 4 kb, 3 kb, 2 kb, 1 kb, 0.5 kb or 0.1 kb of nucleotides which flank the

nucleic acid molecule in the genomic DNA of the cell from which the nucleic acid

molecule is derived.

The nucleic acid molecule can be fused to other coding or regulatory

sequences and still be considered isolated. Thus, recombinant DNA contained in a

20 vector is included in the definition of "isolated" as used herein. Also, isolated

nucleic acid molecules include recombinant DNA molecules in heterologous host

cells, as well as partially or substantially purified DNA molecules in solution.

"Isolated" nucleic acid molecules also encompass *in vivo* and *in vitro* RNA

transcripts of the DNA molecules of the present invention. An isolated nucleic acid

25 molecule or nucleotide sequence can include a nucleic acid molecule or nucleotide

sequence which is synthesized chemically or by recombinant means. Therefore,

recombinant DNA contained in a vector are included in the definition of "isolated"

as used herein. Also, isolated nucleotide sequences include recombinant DNA

molecules in heterologous organisms, as well as partially or substantially purified

30 DNA molecules in solution. *In vivo* and *in vitro* RNA transcripts of the DNA

molecules of the present invention are also encompassed by "isolated" nucleotide

sequences. Such isolated nucleotide sequences are useful in the manufacture of the encoded polypeptide, as probes for isolating homologous sequences (e.g., from other mammalian species), for gene mapping (e.g., by *in situ* hybridization with chromosomes), or for detecting expression of the gene in tissue (e.g., human tissue),

5   such as by Northern blot analysis.

The present invention also pertains to variant nucleic acid molecules which are not necessarily found in nature but which encode a PDE4D polypeptide (e.g., a polypeptide having the amino acid sequence of SEQ ID NO: 2, 3, 4, 5, 6, 7, 8, 9, 10, 12 or 14, or another splicing variant of PDE4D polypeptide or polymorphic variant

10   thereof. Thus, for example, DNA molecules which comprise a sequence that is different from the naturally-occurring nucleotide sequence but which, due to the degeneracy of the genetic code, encode a PDE4D polypeptide of the present invention are also the subject of this invention. The invention also encompasses nucleotide sequences encoding portions (fragments), or encoding variant

15   polypeptides such as analogues or derivatives of the PDE4D polypeptide. Such variants can be naturally-occurring, such as in the case of allelic variation or single nucleotide polymorphisms, or non-naturally-occurring, such as those induced by various mutagens and mutagenic processes. Intended variations include, but are not limited to, addition, deletion and substitution of one or more nucleotides which can

20   result in conservative or non-conservative amino acid changes, including additions and deletions. Preferably the nucleotide (and/or resultant amino acid) changes are silent or conserved; that is, they do not alter the characteristics or activity of the PDE4D polypeptide. In one preferred embodiment, the nucleotide sequences are fragments that comprise one or more polymorphic microsatellite markers. In

25   another preferred embodiment, the nucleotide sequences are fragments that comprise one or more single nucleotide polymorphisms in the PDE4D gene.

Other alterations of the nucleic acid molecules of the invention can include, for example, labeling, methylation, internucleotide modifications such as uncharged linkages (e.g., methyl phosphonates, phosphotriesters, phosphoamidates,

30   carbamates), charged linkages (e.g., phosphorothioates, phosphorodithioates), pendent moieties (e.g., polypeptides), intercalators (e.g., acridine, psoralen),

chelators, alkylators, and modified linkages (e.g., alpha anomeric nucleic acids).
Also included are synthetic molecules that mimic nucleic acid molecules in the
ability to bind to a designated sequences via hydrogen bonding and other chemical
interactions. Such molecules include, for example, those in which peptide linkages

5   substitute for phosphate linkages in the backbone of the molecule.

       The invention also pertains to nucleic acid molecules which hybridize under
high stringency hybridization conditions, such as for selective hybridization, to a
nucleotide sequence described herein (e.g., nucleic acid molecules which
specifically hybridize to a nucleotide sequence encoding polypeptides described

10   herein, and, optionally, have an activity of the polypeptide). In one embodiment, the
invention includes variants described herein which hybridize under high stringency
hybridization conditions (e.g., for selective hybridization) to a nucleotide sequence
comprising a nucleotide sequence selected from SEQ ID NO: 1 which may
optionally comprise at least one polymorphism as shown in Tables 9 and 10 or the

15   complement thereof. In another embodiment, the invention includes variants
described herein which hybridize under high stringency hybridization conditions
(e.g., for selective hybridization) to a nucleotide sequence encoding an amino acid
sequence selected from SEQ ID NO: 2, 3, 4, 5, 6, 7, 8, 9, 10, 12 or 14 or
polymorphic variant thereof. In a preferred embodiment, the variant which

20   hybridizes under high stringency hybridizations has an activity of PDE4D.

       Such nucleic acid molecules can be detected and/or isolated by specific
hybridization (e.g., under high stringency conditions). "Specific hybridization," as
used herein, refers to the ability of a first nucleic acid to hybridize to a second
nucleic acid in a manner such that the first nucleic acid does not hybridize to any

25   nucleic acid other than to the second nucleic acid (e.g., when the first nucleic acid
has a higher similarity to the second nucleic acid than to any other nucleic acid in a
sample wherein the hybridization is to be performed). "Stringency conditions" for
hybridization is a term of art which refers to the incubation and wash conditions,
e.g., conditions of temperature and buffer concentration, which permit hybridization

30   of a particular nucleic acid to a second nucleic acid; the first nucleic acid may be
perfectly (i.e., 100%) complementary to the second, or the first and second may

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share some degree of complementarity which is less than perfect (e.g., 70%, 75%, 85%, 95%). For example, certain high stringency conditions can be used which distinguish perfectly complementary nucleic acids from those of less complementarity. "High stringency conditions", "moderate stringency conditions" and "low stringency conditions" for nucleic acid hybridizations are explained on pages 2.10.1-2.10.16 and pages 6.3.1-6.3.6 in *Current Protocols in Molecular Biology* (Ausubel, F.M. *et al.*, "*Current Protocols in Molecular Biology*", John Wiley & Sons, (1998), the entire teachings of which are incorporated by reference herein). The exact conditions which determine the stringency of hybridization depend not only on ionic strength (e.g., 0.2XSSC, 0.1XSSC), temperature (e.g., room temperature, 42°C, 68°C) and the concentration of destabilizing agents such as formamide or denaturing agents such as SDS, but also on factors such as the length of the nucleic acid sequence, base composition, percent mismatch between hybridizing sequences and the frequency of occurrence of subsets of that sequence within other non-identical sequences. Thus, equivalent conditions can be determined by varying one or more of these parameters while maintaining a similar degree of identity or similarity between the two nucleic acid molecules. Typically, conditions are used such that sequences at least about 60%, at least about 70%, at least about 80%, at least about 90% or at least about 95% or more identical to each other remain hybridized to one another. By varying hybridization conditions from a level of stringency at which no hybridization occurs to a level at which hybridization is first observed, conditions which will allow a given sequence to hybridize (e.g., selectively) with the most similar sequences in the sample can be determined.

Exemplary conditions are described in Krause, M.H. and S.A. Aaronson, *Methods in Enzymology, 200*:546-556 (1991). Also, in, Ausubel, *et al.*, "*Current Protocols in Molecular Biology*", John Wiley & Sons, (1998), which describes the determination of washing conditions for moderate or low stringency conditions. Washing is the step in which conditions are usually set so as to determine a minimum level of complementarity of the hybrids. Generally, starting from the lowest temperature at which only homologous hybridization occurs, each °C by

which the final wash temperature is reduced (holding SSC concentration constant) allows an increase by 1% in the maximum extent of mismatching among the sequences that hybridize. Generally, doubling the concentration of SSC results in an increase in T_m of ~17°C. Using these guidelines, the washing temperature can be

5   determined empirically for high, moderate or low stringency, depending on the level of mismatch sought.

For example, a low stringency wash can comprise washing in a solution containing 0.2XSSC/0.1% SDS for 10 min at room temperature; a moderate stringency wash can comprise washing in a prewarmed solution (42°C) solution

10   containing 0.2XSSC/0.1% SDS for 15 min at 42°C; and a high stringency wash can comprise washing in prewarmed (68°C) solution containing 0.1XSSC/0.1%SDS for 15 min at 68°C. Furthermore, washes can be performed repeatedly or sequentially to obtain a desired result as known in the art. Equivalent conditions can be determined by varying one or more of the parameters given as an example, as known in the art,

15   while maintaining a similar degree of identity or similarity between the target nucleic acid molecule and the primer or probe used.

The percent identity of two nucleotide or amino acid sequences can be determined by aligning the sequences for optimal comparison purposes (*e.g.*, gaps can be introduced in the sequence of a first sequence). The nucleotides or amino

20   acids at corresponding positions are then compared, and the percent identity between the two sequences is a function of the number of identical positions shared by the sequences (*i.e.*, % identity = # of identical positions/total # of positions x 100). In certain embodiments, the length of a sequence aligned for comparison purposes is at least 30%, preferably at least 40%, more preferably at least 60%, and even more

25   preferably at least 70%, 80%, 90% or 95% of the length of the reference sequence. The actual comparison of the two sequences can be accomplished by well-known methods, for example, using a mathematical algorithm. A preferred, non-limiting example of such a mathematical algorithm is described in Karlin *et al.*, *Proc. Natl. Acad. Sci. USA*, *90*:5873-5877 (1993). Such an algorithm is incorporated into the

30   NBLAST and XBLAST programs (version 2.0) as described in Altschul *et al.*, *Nucleic Acids Res.*, *25*:389-3402 (1997). When utilizing BLAST and Gapped

BLAST programs, the default parameters of the respective programs (*e.g.*,
NBLAST) can be used. See http://www.ncbi.nlm.nih.gov. In one embodiment,
parameters for sequence comparison can be set at score=100, wordlength=12, or can
be varied (*e.g.*, W=5 or W=20).

5   Another preferred, non-limiting example of a mathematical algorithm
utilized for the comparison of sequences is the algorithm of Myers and Miller,
CABIOS (1989). Such an algorithm is incorporated into the ALIGN program
(version 2.0) which is part of the GCG sequence alignment software package. When
utilizing the ALIGN program for comparing amino acid sequences, a PAM120

10   weight residue table, a gap length penalty of 12 , and a gap penalty of 4 can be used.
Additional algorithms for sequence analysis are known in the art and include
ADVANCE and ADAM as described in Torellis and Robotti (1994) *Comput. Appl.
Biosci., 10*:3-5; and FASTA described in Pearson and Lipman (1988) *PNAS,
85*:2444-8.

15   In another embodiment, the percent identity between two amino acid
sequences can be accomplished using the GAP program in the CGC software
package (available at http://www.cgc.com) using either a Blossom 63 matrix or a
PAM250 matrix, and a gap weight of 12, 10, 8, 6, or 4 and a length weight of 2, 3, or
4. In yet another embodiment, the percent identity between two nucleic acid

20   sequences can be accomplished using the GAP program in the GCG software
package (available at http://www.accelrys.com), using a gap weight of 50 and a
length weight of 3.

     The present invention also provides isolated nucleic acid molecules that
contain a fragment or portion that hybridizes under highly stringent conditions to a

25   nucleotide sequence comprising a nucleotide sequence selected from SEQ ID NO: 1
which may optionally comprise at least one polymorphism as shown in Tables 9 and
10 and the complement thereof, and also provides isolated nucleic acid molecules
that contain a fragment or portion that hybridizes under highly stringent conditions
to a nucleotide sequence encoding an amino acid sequence selected from SEQ ID

30   NO: 2, 3, 4, 5, 6, 7, 8, 9, 10, 12 or 14, or polymorphic variant thereof. The nucleic
acid fragments of the invention are at least about 15, preferably at least about 18, 20,

23 or 25 nucleotides, and can be 30, 40, 50, 100, 200 or more nucleotides in length.
Longer fragments, for example, 30 or more nucleotides in length, which encode
antigenic polypeptides described herein are particularly useful, such as for the
generation of antibodies as described below.

5   In a related aspect, the nucleic acid fragments of the invention are used as
probes or primers in assays such as those described herein. "Probes" or "primers"
are oligonucleotides that hybridize in a base-specific manner to a complementary
strand of nucleic acid molecules. Such probes and primers include polypeptide
nucleic acids, as described in Nielsen *et al.*, *Science, 254*, 1497-1500 (1991).

10   Typically, a probe or primer comprises a region of nucleotide sequence that
hybridizes to at least about 15, typically about 20-25, and more typically about 40,
50 or 75, consecutive nucleotides of a nucleic acid molecule comprising a
contiguous nucleotide sequence selected from: SEQ ID NO: 1 which may optionally
comprise at least one polymorphism as shown in Tables 9 and 10, the complement

15   thereof, or a sequence encoding an amino acid sequence selected from SEQ ID NO:
2, 3, 4, 5, 6, 7, 8, 9, 10, 12 or 14, or polymorphic variant thereof. In preferred
embodiments, a probe or primer comprises 100 or fewer nucleotides, preferably
from 6 to 50 nucleotides, preferably from 12 to 30 nucleotides. In other
embodiments, the probe or primer is at least 70% identical to the contiguous

20   nucleotide sequence or to the complement of the contiguous nucleotide sequence,
preferably at least 80% identical, more preferably at least 90% identical, even more
preferably at least 95% identical, or even capable of selectively hybridizing to the
contiguous nucleotide sequence or to the complement of the contiguous nucleotide
sequence. Often, the probe or primer further comprises a label, *e.g.*, radioisotope,

25   fluorescent compound, enzyme, or enzyme co-factor.

           The nucleic acid molecules of the invention such as those described above
can be identified and isolated using standard molecular biology techniques and the
sequence information provided herein. For example, nucleic acid molecules can be
amplified and isolated by the polymerase chain reaction using synthetic

30   oligonucleotide primers designed based on one or more of the sequences provided in
SEQ ID NO: 1 which may optionally comprise at least one polymorphism shown in

Tables 9 and 10, and/or the complement thereof, or designed on nucleotides based on sequences encoding one or more of the amino acid sequences provided herein. See generally *PCR Technology: Principles and Applications for DNA Amplification* (ed. H.A. Erlich, Freeman Press, NY, NY, 1992); *PCR Protocols: A*

5   *Guide to Methods and Applications* (Eds. Innis, *et al.*, Academic Press, San Diego, CA, 1990); Mattila *et al.*, *Nucleic Acids Res., 19*:4967 (1991); Eckert *et al.*, *PCR Methods and Applications, 1*:17 (1991); PCR (eds. McPherson *et al.*, IRL Press, Oxford); and U.S. Patent 4,683,202. The nucleic acid molecules can be amplified using cDNA, mRNA or genomic DNA as a template, cloned into an appropriate

10   vector and characterized by DNA sequence analysis.

Other suitable amplification methods include the ligase chain reaction (LCR) (see Wu and Wallace, *Genomics, 4*:560 (1989), Landegren *et al.*, *Science, 241*:1077 (1988), transcription amplification (Kwoh *et al.*, *Proc. Natl. Acad. Sci. USA, 86*:1173 (1989)), and self-sustained sequence replication (Guatelli *et al.*, *Proc. Nat.*

15   *Acad. Sci. USA, 87*:1874 (1990)) and nucleic acid based sequence amplification (NASBA). The latter two amplification methods involve isothermal reactions based on isothermal transcription, which produce both single stranded RNA (ssRNA) and double stranded DNA (dsDNA) as the amplification products in a ratio of about 30 or 100 to 1, respectively.

20   The amplified DNA can be radiolabelled and used as a probe for screening a cDNA library derived from human cells, mRNA in zap express, ZIPLOX or other suitable vector. Corresponding clones can be isolated, DNA can obtained following *in vivo* excision, and the cloned insert can be sequenced in either or both orientations by art recognized methods to identify the correct reading frame encoding a

25   polypeptide of the appropriate molecular weight. For example, the direct analysis of the nucleotide sequence of nucleic acid molecules of the present invention can be accomplished using well-known methods that are commercially available. See, for example, Sambrook *et al.*, *Molecular Cloning, A Laboratory Manual* (2nd Ed., CSHP, New York 1989); Zyskind *et al.*, *Recombinant DNA Laboratory Manual*,

30   (Acad. Press, 1988)). Using these or similar methods, the polypeptide and the DNA encoding the polypeptide can be isolated, sequenced and further characterized.

Antisense nucleic acid molecules of the invention can be designed using the nucleotide sequences of SEQ ID NO: 1 and/or the complement of SEQ ID NO: 1, and/or a portion of SEQ ID NO:1 or the complement of SEQ ID NO:1 and/or a sequence encoding the amino acid sequences or SEQ ID NO: 2, 3, 4, 5, 6, 7, 8, 9, 10,

5   12 and/or 14, or encoding a portion of SEQ ID NO: 2, 3, 4, 5, 6, 7, 8, 9, 10, 12 and/or 14, (wherein any one of these may optionally comprise at least one polymorphism as shown in Tables 9 and 10) and constructed using chemical synthesis and enzymatic ligation reactions using procedures known in the art. For example, an antisense nucleic acid molecule (*e.g.*, an antisense oligonucleotide) can

10   be chemically synthesized using naturally occurring nucleotides or variously modified nucleotides designed to increase the biological stability of the molecules or to increase the physical stability of the duplex formed between the antisense and sense nucleic acids, *e.g.*, phosphorothioate derivatives and acridine substituted nucleotides can be used. Alternatively, the antisense nucleic acid molecule can be

15   produced biologically using an expression vector into which a nucleic acid molecule has been subcloned in an antisense orientation (*i.e.*, RNA transcribed from the inserted nucleic acid molecule will be of an antisense orientation to a target nucleic acid of interest).

In general, the isolated nucleic acid sequences of the invention can be used as

20   molecular weight markers on Southern gels, and as chromosome markers which are labeled to map related gene positions. The nucleic acid sequences can also be used to compare with endogenous DNA sequences in patients to identify genetic disorders (e.g., a predisposition for or susceptibility to stroke), and as probes, such as to hybridize and discover related DNA sequences or to subtract out known

25   sequences from a sample. The nucleic acid sequences can further be used to derive primers for genetic fingerprinting, to raise anti-polypeptide antibodies using DNA immunization techniques, and as an antigen to raise anti-DNA antibodies or elicit immune responses. Portions or fragments of the nucleotide sequences identified herein (and the corresponding complete gene sequences) can be used in numerous

30   ways as polynucleotide reagents. For example, these sequences can be used to: (i) map their respective genes on a chromosome; and, thus, locate gene regions

associated with genetic disease; (ii) identify an individual from a minute biological
sample (tissue typing); and (iii) aid in forensic identification of a biological sample.
Additionally, the nucleotide sequences of the invention can be used to identify and
express recombinant polypeptides for analysis, characterization or therapeutic use, or

5   as markers for tissues in which the corresponding polypeptide is expressed, either
constitutively, during tissue differentiation, or in diseased states. The nucleic acid
sequences can additionally be used as reagents in the screening and/or diagnostic
assays described herein, and can also be included as components of kits (e.g.,
reagent kits) for use in the screening and/or diagnostic assays described herein.

10   Another aspect of the invention pertains to nucleic acid constructs containing
a nucleic acid molecule selected from the group consisting of SEQ ID NO: 1 which
may optionally comprise at least one polymorphism shown in Tables 9 and 10 and
the complement thereof (or a portion thereof). Yet another aspect of the invention
pertains to nucleic acid constructs containing a nucleic acid molecule encoding the

15   amino acid sequence of SEQ ID NO: 2, 3, 4, 5, 6, 7, 8, 9, 10, 12 or 14 or
polymorphic variant thereof. The constructs comprise a vector (e.g., an expression
vector) into which a sequence of the invention has been inserted in a sense or
antisense orientation. As used herein, the term "vector" refers to a nucleic acid
molecule capable of transporting another nucleic acid to which it has been linked.

20   One type of vector is a "plasmid", which refers to a circular double stranded DNA
loop into which additional DNA segments can be ligated. Another type of vector is
a viral vector, wherein additional DNA segments can be ligated into the viral
genome. Certain vectors are capable of autonomous replication in a host cell into
which they are introduced (e.g., bacterial vectors having a bacterial origin of

25   replication and episomal mammalian vectors). Other vectors (e.g., non-episomal
mammalian vectors) are integrated into the genome of a host cell upon introduction
into the host cell, and thereby are replicated along with the host genome. Moreover,
certain vectors, expression vectors, are capable of directing the expression of genes
to which they are operably linked. In general, expression vectors of utility in

30   recombinant DNA techniques are often in the form of plasmids. However, the
invention is intended to include such other forms of expression vectors, such as viral

vectors (*e.g.*, replication defective retroviruses, adenoviruses and adeno-associated viruses) that serve equivalent functions.

Preferred recombinant expression vectors of the invention comprise a nucleic acid molecule of the invention in a form suitable for expression of the nucleic acid

5   molecule in a host cell. This means that the recombinant expression vectors include one or more regulatory sequences, selected on the basis of the host cells to be used for expression, which is operably linked to the nucleic acid sequence to be expressed. Within a recombinant expression vector, "operably or operatively linked" is intended to mean that the nucleotide sequence of interest is linked to the

10   regulatory sequence(s) in a manner which allows for expression of the nucleotide sequence (*e.g.*, in an *in vitro* transcription/translation system or in a host cell when the vector is introduced into the host cell). The term "regulatory sequence" is intended to include promoters, enhancers and other expression control elements (*e.g.*, polyadenylation signals). Such regulatory sequences are described, for

15   example, in Goeddel, *Gene Expression Technology: Methods in Enzymology 185*, Academic Press, San Diego, CA (1990). Regulatory sequences include those which direct constitutive expression of a nucleotide sequence in many types of host cell and those which direct expression of the nucleotide sequence only in certain host cells (*e.g.*, tissue-specific regulatory sequences). It will be appreciated by those

20   skilled in the art that the design of the expression vector can depend on such factors as the choice of the host cell to be transformed and the level of expression of polypeptide desired. The expression vectors of the invention can be introduced into host cells to thereby produce polypeptides, including fusion polypeptides, encoded by nucleic acid molecules as described herein.

25   The recombinant expression vectors of the invention can be designed for expression of a polypeptide of the invention in prokaryotic or eukaryotic cells, *e.g.*, bacterial cells such as *E. coli*, insect cells (using baculovirus expression vectors), yeast cells or mammalian cells. Suitable host cells are discussed further in Goeddel, *supra*. Alternatively, the recombinant expression vector can be transcribed and

30   translated *in vitro*, for example using T7 promoter regulatory sequences and T7 polymerase.

Another aspect of the invention pertains to host cells into which a recombinant expression vector of the invention has been introduced. The terms "host cell" and "recombinant host cell" are used interchangeably herein. It is understood that such terms refer not only to the particular subject cell but also to the

5   progeny or potential progeny of such a cell. Because certain modifications may occur in succeeding generations due to either mutation or environmental influences, such progeny may not, in fact, be identical to the parent cell, but are still included within the scope of the term as used herein.

A host cell can be any prokaryotic or eukaryotic cell. For example, a nucleic

10   acid molecule of the invention can be expressed in bacterial cells (*e.g.*, *E. coli*), insect cells, yeast or mammalian cells (such as Chinese hamster ovary cells (CHO) or COS cells). Other suitable host cells are known to those skilled in the art.

Vector DNA can be introduced into prokaryotic or eukaryotic cells via conventional transformation or transfection techniques. As used herein, the terms

15   "transformation" and "transfection" are intended to refer to a variety of art-recognized techniques for introducing a foreign nucleic acid molecule (*e.g.*, DNA) into a host cell, including calcium phosphate or calcium chloride co-precipitation, DEAE-dextran-mediated transfection, lipofection, or electroporation. Suitable methods for transforming or transfecting host cells can be

20   found in Sambrook, *et al.* (*supra*), and other laboratory manuals.

For stable transfection of mammalian cells, it is known that, depending upon the expression vector and transfection technique used, only a small fraction of cells may integrate the foreign DNA into their genome. In order to identify and select these integrants, a gene that encodes a selectable marker (*e.g.*, for resistance to

25   antibiotics) is generally introduced into the host cells along with the gene of interest. Preferred selectable markers include those that confer resistance to drugs, such as G418, hygromycin and methotrexate. Nucleic acid molecules encoding a selectable marker can be introduced into a host cell on the same vector as the nucleic acid molecule of the invention or can be introduced on a separate vector. Cells stably

30   transfected with the introduced nucleic acid molecule can be identified by drug

selection (*e.g.*, cells that have incorporated the selectable marker gene will survive, while the other cells die).

A host cell of the invention, such as a prokaryotic or eukaryotic host cell in culture, can be used to produce (*i.e.*, express) a polypeptide of the invention.

5   Accordingly, the invention further provides methods for producing a polypeptide using the host cells of the invention. In one embodiment, the method comprises culturing the host cell of invention (into which a recombinant expression vector encoding a polypeptide of the invention has been introduced) in a suitable medium such that the polypeptide is produced. In another embodiment, the method further

10   comprises isolating the polypeptide from the medium or the host cell.

The host cells of the invention can also be used to produce nonhuman transgenic animals. For example, in one embodiment, a host cell of the invention is a fertilized oocyte or an embryonic stem cell into which a nucleic acid molecule of the invention has been introduced (e.g., an exogenous PDE4D gene, or an exogenous

15   nucleic acid encoding PDE4D polypeptide). Such host cells can then be used to create non-human transgenic animals in which exogenous nucleotide sequences have been introduced into the genome or homologous recombinant animals in which endogenous nucleotide sequences have been altered. Such animals are useful for studying the function and/or activity of the nucleotide sequence and polypeptide

20   encoded by the sequence and for identifying and/or evaluating modulators of their activity. As used herein, a "transgenic animal" is a non-human animal, preferably a mammal, more preferably a rodent such as a rat or mouse, in which one or more of the cells of the animal includes a transgene. Other examples of transgenic animals include non-human primates, sheep, dogs, cows, goats, chickens and amphibians. A

25   transgene is exogenous DNA which is integrated into the genome of a cell from which a transgenic animal develops and which remains in the genome of the mature animal, thereby directing the expression of an encoded gene product in one or more cell types or tissues of the transgenic animal. As used herein, an "homologous recombinant animal" is a non-human animal, preferably a mammal, more preferably

30   a mouse, in which an endogenous gene has been altered by homologous recombination between the endogenous gene and an exogenous DNA molecule

introduced into a cell of the animal, *e.g.*, an embryonic cell of the animal, prior to development of the animal.

Methods for generating transgenic animals via embryo manipulation and microinjection, particularly animals such as mice, have become conventional in the

5   art and are described, for example, in U.S. Patent Nos. 4,736,866 and 4,870,009, U.S. Patent No. 4,873,191 and in Hogan, *Manipulating the Mouse Embryo* (Cold Spring Harbor Laboratory Press, Cold Spring Harbor, N.Y., 1986). Methods for constructing homologous recombination vectors and homologous recombinant animals are described further in Bradley (1991) *Current Opinion in Bio/Technology*,

10   *2*:823-829 and in PCT Publication Nos. WO 90/11354, WO 91/01140, WO 92/0968, and WO 93/04169. Clones of the non-human transgenic animals described herein can also be produced according to the methods described in Wilmut *et al.* (1997) *Nature, 385*:810-813 and PCT Publication Nos. WO 97/07668 and WO 97/07669.

POLYPEPTIDES OF THE INVENTION

15   The present invention also pertains to isolated polypeptides encoded by PDE4D ("PDE4D polypeptides") and fragments and variants thereof, as well as polypeptides encoded by nucleotide sequences described herein (e.g., other splicing variants). The term "polypeptide" refers to a polymer of amino acids, and not to a specific length; thus, peptides, oligopeptides and proteins are included within the

20   definition of a polypeptide. As used herein, a polypeptide is said to be "isolated" or "purified" when it is substantially free of cellular material when it is isolated from recombinant and non-recombinant cells, or free of chemical precursors or other chemicals when it is chemically synthesized. A polypeptide, however, can be joined to another polypeptide with which it is not normally associated in a cell (e.g., in a

25   "fusion protein") and still be "isolated" or "purified."

The polypeptides of the invention can be purified to homogeneity. It is understood, however, that preparations in which the polypeptide is not purified to homogeneity are useful. The critical feature is that the preparation allows for the desired function of the polypeptide, even in the presence of considerable amounts of

30   other components. Thus, the invention encompasses various degrees of purity. In

one embodiment, the language "substantially free of cellular material" includes preparations of the polypeptide having less than about 30% (by dry weight) other proteins (*i.e.*, contaminating protein), less than about 20% other proteins, less than about 10% other proteins, or less than about 5% other proteins.

5   When a polypeptide is recombinantly produced, it can also be substantially free of culture medium, *i.e.*, culture medium represents less than about 20%, less than about 10%, or less than about 5% of the volume of the polypeptide preparation. The language "substantially free of chemical precursors or other chemicals" includes preparations of the polypeptide in which it is separated from chemical precursors or

10   other chemicals that are involved in its synthesis. In one embodiment, the language "substantially free of chemical precursors or other chemicals" includes preparations of the polypeptide having less than about 30% (by dry weight) chemical precursors or other chemicals, less than about 20% chemical precursors or other chemicals, less than about 10% chemical precursors or other chemicals, or less than about 5%

15   chemical precursors or other chemicals.

In one embodiment, a polypeptide of the invention comprises an amino acid sequence encoded by a nucleic acid molecule comprising a nucleotide sequence selected from the group consisting of SEQ ID NO: 1 which may optionally comprise at least one polymorphism shown in Tables 9 and 10 and complements and portions

20   thereof, e.g., SEQ ID NO: 2, 3, 4, 5, 6, 7, 8, 9, 10, 12 or 14, or a portion or polymorphic variant thereof. However, the polypeptides of the invention also encompass fragment and sequence variants. Variants include a substantially homologous polypeptide encoded by the same genetic locus in an organism, *i.e.*, an allelic variant, as well as other splicing variants. Variants also encompass

25   polypeptides derived from other genetic loci in an organism, but having substantial homology to a polypeptide encoded by a nucleic acid molecule comprising a nucleotide sequence selected from the group consisting of SEQ ID NO: 1 which may optionally comprise at least one polymorphism shown in Tables 9 and 10 and complements and portions thereof, or having substantial homology to a polypeptide

30   encoded by a nucleic acid molecule comprising a nucleotide sequence selected from the group consisting of nucleotide sequences encoding SEQ ID NO: 2, 3, 4, 5, 6, 7,

8, 9, 10, 12 or 14, or polymorphic variants thereof. Variants also include polypeptides substantially homologous or identical to these polypeptides but derived from another organism, *i.e.*, an ortholog. Variants also include polypeptides that are substantially homologous or identical to these polypeptides that are produced by

5   chemical synthesis. Variants also include polypeptides that are substantially homologous or identical to these polypeptides that are produced by recombinant methods.

As used herein, two polypeptides (or a region of the polypeptides) are substantially homologous or identical when the amino acid sequences are at least

10   about 45-55%, typically at least about 70-75%, more typically at least about 80-85%, and most typically greater than about 90% or more homologous or identical. A substantially homologous amino acid sequence, according to the present invention, will be encoded by a nucleic acid molecule hybridizing to SEQ ID NO: 1 which may optionally comprise at least one polymorphism shown in Tables 9 and 10, or portion

15   thereof, under stringent conditions as more particularly described above, or will be encoded by a nucleic acid molecule hybridizing to a nucleic acid sequence encoding SEQ ID NO: 2, 3, 4, 5, 6, 7, 8, 9, 10, 12 or 14, portion thereof or polymorphic variant thereof, under stringent conditions as more particularly described thereof.

To determine the percent homology or identity of two amino acid sequences,

20   or of two nucleic acid sequences, the sequences are aligned for optimal comparison purposes (*e.g.*, gaps can be introduced in the sequence of one polypeptide or nucleic acid molecule for optimal alignment with the other polypeptide or nucleic acid molecule). The amino acid residues or nucleotides at corresponding amino acid positions or nucleotide positions are then compared. When a position in one

25   sequence is occupied by the same amino acid residue or nucleotide as the corresponding position in the other sequence, then the molecules are homologous at that position. As used herein, amino acid or nucleic acid "homology" is equivalent to amino acid or nucleic acid "identity". The percent homology between the two sequences is a function of the number of identical positions shared by the sequences

30   (*i.e.*, percent homology equals the number of identical positions/total number of positions times 100).

The invention also encompasses polypeptides having a lower degree of identity but having sufficient similarity so as to perform one or more of the same functions performed by a polypeptide encoded by a nucleic acid molecule of the invention. Similarity is determined by conserved amino acid substitution. Such

5   substitutions are those that substitute a given amino acid in a polypeptide by another amino acid of like characteristics. Conservative substitutions are likely to be phenotypically silent. Typically seen as conservative substitutions are the replacements, one for another, among the aliphatic amino acids Ala, Val, Leu and Ile; interchange of the hydroxyl residues Ser and Thr, exchange of the acidic

10 residues Asp and Glu, substitution between the amide residues Asn and Gln, exchange of the basic residues Lys and Arg and replacements among the aromatic residues Phe and Tyr. Guidance concerning which amino acid changes are likely to be phenotypically silent are found in Bowie *et al.*, *Science 247*:1306-1310 (1990).

A variant polypeptide can differ in amino acid sequence by one or more

15 substitutions, deletions, insertions, inversions, fusions, and truncations or a combination of any of these. Further, variant polypeptides can be fully functional or can lack function in one or more activities. Fully functional variants typically contain only conservative variation or variation in non-critical residues or in non-critical regions. Functional variants can also contain substitution of similar

20 amino acids that result in no change or an insignificant change in function. Alternatively, such substitutions may positively or negatively affect function to some degree. Non-functional variants typically contain one or more non-conservative amino acid substitutions, deletions, insertions, inversions, or truncation or a substitution, insertion, inversion, or deletion in a critical residue or critical region.

25 Amino acids that are essential for function can be identified by methods known in the art, such as site-directed mutagenesis or alanine-scanning mutagenesis (Cunningham *et al.*, *Science, 244*:1081-1085 (1989)). The latter procedure introduces single alanine mutations at every residue in the molecule. The resulting mutant molecules are then tested for biological activity *in vitro*, or *in vitro*

30 proliferative activity. Sites that are critical for polypeptide activity can also be determined by structural analysis such as crystallization, nuclear magnetic resonance

or photoaffinity labeling (Smith *et al.*, *J. Mol. Biol.*, 224:899-904 (1992); de Vos *et al.*, *Science*, 255:306-312 (1992)).

The invention also includes polypeptide fragments of the polypeptides of the invention. Fragments can be derived from a polypeptide encoded by a nucleic acid
5 molecule comprising SEQ ID NO: 1 which may optionally comprise at least one polymorphism shown in Tables 9 and 10 or a portion thereof and the complements thereof (e.g., SEQ ID NO: 2, 3, 4, 5, 6, 7, 8, 9, 10, 12 or 14, or other splicing variants). However, the invention also encompasses fragments of the variants of the polypeptides described herein. As used herein, a fragment comprises at least 6
10 contiguous amino acids. Useful fragments include those that retain one or more of the biological activities of the polypeptide as well as fragments that can be used as an immunogen to generate polypeptide-specific antibodies.

Biologically active fragments (peptides which are, for example, 6, 9, 12, 15, 16, 20, 30, 35, 36, 37, 38, 39, 40, 50, 100 or more amino acids in length) can
15 comprise a domain, segment, or motif that has been identified by analysis of the polypeptide sequence using well-known methods, *e.g.*, signal peptides, extracellular domains, one or more transmembrane segments or loops, ligand binding regions, zinc finger domains, DNA binding domains, acylation sites, glycosylation sites, or phosphorylation sites.

20 Fragments can be discrete (not fused to other amino acids or polypeptides) or can be within a larger polypeptide. Further, several fragments can be comprised within a single larger polypeptide. In one embodiment a fragment designed for expression in a host can have heterologous pre- and pro-polypeptide regions fused to the amino terminus of the polypeptide fragment and an additional region fused to the
25 carboxyl terminus of the fragment.

The invention thus provides chimeric or fusion polypeptides. These comprise a polypeptide of the invention operatively linked to a heterologous protein or polypeptide having an amino acid sequence not substantially homologous to the polypeptide. "Operatively linked" indicates that the polypeptide and the
30 heterologous protein are fused in-frame. The heterologous protein can be fused to the N-terminus or C-terminus of the polypeptide. In one embodiment the fusion

polypeptide does not affect function of the polypeptide *per se*. For example, the fusion polypeptide can be a GST-fusion polypeptide in which the polypeptide sequences are fused to the C-terminus of the GST sequences. Other types of fusion polypeptides include, but are not limited to, enzymatic fusion polypeptides, for
5   example β-galactosidase fusions, yeast two-hybrid GAL fusions, poly-His fusions and Ig fusions. Such fusion polypeptides, particularly poly-His fusions, can facilitate the purification of recombinant polypeptide. In certain host cells (*e.g.*, mammalian host cells), expression and/or secretion of a polypeptide can be increased by using a heterologous signal sequence. Therefore, in another
10   embodiment, the fusion polypeptide contains a heterologous signal sequence at its N-terminus.

EP-A-O 464 533 discloses fusion proteins comprising various portions of immunoglobulin constant regions. The Fc is useful in therapy and diagnosis and thus results, for example, in improved pharmacokinetic properties (EP-A 0232 262).
15   In drug discovery, for example, human proteins have been fused with Fc portions for the purpose of high-throughput screening assays to identify antagonists. Bennett *et al.*, *Journal of Molecular Recognition,* 8:52-58 (1995) and Johanson *et al.*, *The Journal of Biological Chemistry,* 270,16:9459-9471 (1995). Thus, this invention also encompasses soluble fusion polypeptides containing a polypeptide of the
20   invention and various portions of the constant regions of heavy or light chains of immunoglobulins of various subclass (IgG, IgM, IgA, IgE).

A chimeric or fusion polypeptide can be produced by standard recombinant DNA techniques. For example, DNA fragments coding for the different polypeptide sequences are ligated together in-frame in accordance with conventional techniques.
25   In another embodiment, the fusion gene can be synthesized by conventional techniques including automated DNA synthesizers. Alternatively, PCR amplification of nucleic acid fragments can be carried out using anchor primers which give rise to complementary overhangs between two consecutive nucleic acid fragments which can subsequently be annealed and re-amplified to generate a chimeric nucleic acid
30   sequence (see Ausubel *et al.*, *Current Protocols in Molecular Biology,* 1992). Moreover, many expression vectors are commercially available that already encode a

fusion moiety (*e.g.*, a GST protein). A nucleic acid molecule encoding a polypeptide of the invention can be cloned into such an expression vector such that the fusion moiety is linked in-frame to the polypeptide.

The isolated polypeptide can be purified from cells that naturally express it,
5   purified from cells that have been altered to express it (recombinant), or synthesized using known protein synthesis methods. In one embodiment, the polypeptide is produced by recombinant DNA techniques. For example, a nucleic acid molecule encoding the polypeptide is cloned into an expression vector, the expression vector introduced into a host cell and the polypeptide expressed in the host cell. The
10   polypeptide can then be isolated from the cells by an appropriate purification scheme using standard protein purification techniques.

In general, polypeptides of the present invention can be used as a molecular weight marker on SDS-PAGE gels or on molecular sieve gel filtration columns using art-recognized methods. The polypeptides of the present invention can be
15   used to raise antibodies or to elicit an immune response. The polypeptides can also be used as a reagent, *e.g.*, a labeled reagent, in assays to quantitatively determine levels of the polypeptide or a molecule to which it binds (*e.g.*, a receptor or a ligand) in biological fluids. The polypeptides can also be used as markers for cells or tissues in which the corresponding polypeptide is preferentially expressed, either
20   constitutively, during tissue differentiation, or in a diseased state. The polypeptides can be used to isolate a corresponding binding agent, *e.g.*, receptor or ligand, such as, for example, in an interaction trap assay, and to screen for peptide or small molecule antagonists or agonists of the binding interaction.

ANTIBODIES OF THE INVENTION
25   Polyclonal and/or monoclonal antibodies that specifically bind one form of the gene product but not to the other form of the gene product are also provided. Antibodies are also provided that bind a portion of either the variant or the reference gene product that contains the polymorphic site or sites. The invention provides antibodies to the polypeptides and polypeptide fragments of the invention, *e.g.*,
30   having an amino acid sequence encoded by SEQ ID NO: 2, 3, 4, 5, 6, 7, 8, 9, 10, 12

or 14, or a portion thereof, or having an amino acid sequence encoded by a nucleic

acid molecule comprising all or a portion of SEQ ID NO: 1 which may optionally

comprise at least one polymorphism shown in Tables 9 and 10 (e.g., SEQ ID NO: 2,

3, 4, 5, 6, 7, 8, 9, 10, 12 or 14, or another splicing variant or portion thereof). The

5   term "antibody" as used herein refers to immunoglobulin molecules and

immunologically active portions of immunoglobulin molecules, *i.e.*, molecules that

contain an antigen binding site that specifically binds an antigen. A molecule that

specifically binds to a polypeptide of the invention is a molecule that binds to that

polypeptide or a fragment thereof, but does not substantially bind other molecules in

10   a sample, *e.g.*, a biological sample, which naturally contains the polypeptide.

Examples of immunologically active portions of immunoglobulin molecules include

F(ab) and F(ab')$_2$ fragments which can be generated by treating the antibody with an

enzyme such as pepsin. The invention provides polyclonal and monoclonal

antibodies that bind to a polypeptide of the invention. The term "monoclonal

15   antibody" or "monoclonal antibody composition", as used herein, refers to a

population of antibody molecules that contain only one species of an antigen binding

site capable of immunoreacting with a particular epitope of a polypeptide of the

invention. A monoclonal antibody composition thus typically displays a single

binding affinity for a particular polypeptide of the invention with which it

20   immunoreacts.

Polyclonal antibodies can be prepared as described above by immunizing a

suitable subject with a desired immunogen, *e.g.*, polypeptide of the invention or

fragment thereof. The antibody titer in the immunized subject can be monitored

over time by standard techniques, such as with an enzyme linked immunosorbent

25   assay (ELISA) using immobilized polypeptide. If desired, the antibody molecules

directed against the polypeptide can be isolated from the mammal (*e.g.*, from the

blood) and further purified by well-known techniques, such as protein A

chromatography to obtain the IgG fraction. At an appropriate time after

immunization, *e.g.*, when the antibody titers are highest, antibody-producing cells

30   can be obtained from the subject and used to prepare monoclonal antibodies by

standard techniques, such as the hybridoma technique originally described by Kohler

-30-

and Milstein (1975) *Nature, 256*:495-497, the human B cell hybridoma technique
(Kozbor *et al.* (1983) *Immunol. Today, 4*:72), the EBV-hybridoma technique (Cole
et al. (1985), *Monoclonal Antibodies and Cancer Therapy*, Alan R. Liss, Inc., pp.
77-96) or trioma techniques. The technology for producing hybridomas is well

5   known (see generally *Current Protocols in Immunology* (1994) Coligan *et al.* (eds.)
John Wiley & Sons, Inc., New York, NY). Briefly, an immortal cell line (typically a
myeloma) is fused to lymphocytes (typically splenocytes) from a mammal
immunized with an immunogen as described above, and the culture supernatants of
the resulting hybridoma cells are screened to identify a hybridoma producing a

10   monoclonal antibody that binds a polypeptide of the invention.

Any of the many well known protocols used for fusing lymphocytes and
immortalized cell lines can be applied for the purpose of generating a monoclonal
antibody to a polypeptide of the invention (see, *e.g., Current Protocols in
Immunology, supra*; Galfre *et al.* (1977) *Nature, 266*:55052; R.H. Kenneth, in

15   *Monoclonal Antibodies: A New Dimension In Biological Analyses*, Plenum
Publishing Corp., New York, New York (1980); and Lerner (1981) *Yale J. Biol.
Med., 54*:387-402. Moreover, the ordinarily skilled worker will appreciate that there
are many variations of such methods that also would be useful.

Alternative to preparing monoclonal antibody-secreting hybridomas, a

20   monoclonal antibody to a polypeptide of the invention can be identified and isolated
by screening a recombinant combinatorial immunoglobulin library (*e.g.*, an antibody
phage display library) with the polypeptide to thereby isolate immunoglobulin
library members that bind the polypeptide. Kits for generating and screening phage
display libraries are commercially available (*e.g.*, the Pharmacia *Recombinant Phage*

25   *Antibody System*, Catalog No. 27-9400-01; and the Stratagene *SurfZAP*™ Phage
Display Kit, Catalog No. 240612). Additionally, examples of methods and reagents
particularly amenable for use in generating and screening antibody display library
can be found in, for example, U.S. Patent No. 5,223,409; PCT Publication No. WO
92/18619; PCT Publication No. WO 91/17271; PCT Publication No. WO 92/20791;

30   PCT Publication No. WO 92/15679; PCT Publication No. WO 93/01288; PCT
Publication No. WO 92/01047; PCT Publication No. WO 92/09690; PCT

Publication No. WO 90/02809; Fuchs *et al.* (1991) *Bio/Technology, 9*:1370-1372; Hay *et al.* (1992) *Hum. Antibod. Hybridomas, 3*:81-85; Huse *et al.* (1989) *Science, 246*:1275-1281; Griffiths *et al.* (1993) *EMBO J., 12*:725-734.

5   Additionally, recombinant antibodies, such as chimeric and humanized monoclonal antibodies, comprising both human and non-human portions, which can be made using standard recombinant DNA techniques, are within the scope of the invention. Such chimeric and humanized monoclonal antibodies can be produced by recombinant DNA techniques known in the art.

In general, antibodies of the invention (*e.g.*, a monoclonal antibody) can be
10   used to isolate a polypeptide of the invention by standard techniques, such as affinity chromatography or immunoprecipitation. A polypeptide-specific antibody can facilitate the purification of natural polypeptide from cells and of recombinantly produced polypeptide expressed in host cells. Moreover, an antibody specific for a polypeptide of the invention can be used to detect the polypeptide (*e.g.*, in a cellular
15   lysate, cell supernatant, or tissue sample) in order to evaluate the abundance and pattern of expression of the polypeptide. Antibodies can be used diagnostically to monitor protein levels in tissue as part of a clinical testing procedure, *e.g.*, to, for example, determine the efficacy of a given treatment regimen. Detection can be facilitated by coupling the antibody to a detectable substance. Examples of
20   detectable substances include various enzymes, prosthetic groups, fluorescent materials, luminescent materials, bioluminescent materials, and radioactive materials. Examples of suitable enzymes include horseradish peroxidase, alkaline phosphatase, β-galactosidase, or acetylcholinesterase; examples of suitable prosthetic group complexes include streptavidin/biotin and avidin/biotin; examples
25   of suitable fluorescent materials include umbelliferone, fluorescein, fluorescein isothiocyanate, rhodamine, dichlorotriazinylamine fluorescein, dansyl chloride or phycoerythrin; an example of a luminescent material includes luminol; examples of bioluminescent materials include luciferase, luciferin, and aequorin, and examples of suitable radioactive material include 125I, 131I, 35S or 3H.

DIAGNOSTIC AND SCREENING ASSAYS OF THE INVENTION

The present invention also pertains to a method of diagnosing or aiding in the diagnosis of stroke associated with the presence of the PDE4D gene or gene product in an individual. Diagnostic assays can be designed for assessing PDE4D gene expression, or for assessing activity of PDE4D polypeptides of the invention. In one embodiment, the assays are used in the context of a biological sample (e.g., blood, serum, cells, tissue) to thereby determine whether an individual is afflicted with stroke, or is at risk for (has a predisposition for or a susceptibility to) developing stroke. The invention also provides for prognostic (or predictive) assays for determining whether an individual is susceptible to developing stroke. For example, mutations in the gene can be assayed in a biological sample. Such assays can be used for prognostic or predictive purpose to thereby prophylactically treat an individual prior to the onset of symptoms associated with stroke. Another aspect of the invention pertains to assays for monitoring the influence of agents (e.g., drugs, compounds or other agents) on the gene expression or activity of polypeptides of the invention, as well as to assays for identifying agents which bind to PDE4D polypeptides. These and other assays and agents are described in further detail in the following sections.

DIAGNOSTIC ASSAYS

The nucleic acids, probes, primers, polypeptides and antibodies described herein can be used in methods of diagnosis of a susceptibility to stroke, as well as in kits useful for diagnosis of a susceptibility to stroke.

In one embodiment of the invention, diagnosis of a susceptibility to stroke is made by detecting a polymorphism in PDE4D as described herein. The polymorphism can be a mutation in PDE4D, such as the insertion or deletion of a single nucleotide, or of more than one nucleotide, resulting in a frame shift mutation; the change of at least one nucleotide, resulting in a change in the encoded amino acid; the change of at least one nucleotide, resulting in the generation of a premature stop codon; the deletion of several nucleotides, resulting in a deletion of one or more amino acids encoded by the nucleotides; the insertion of one or several nucleotides,

such as by unequal recombination or gene conversion, resulting in an interruption of the coding sequence of the gene; duplication of all or a part of the gene; transposition of all or a part of the gene; or rearrangement of all or a part of the gene. More than one such mutation may be present in a single gene. Such sequence

5   changes cause a mutation in the polypeptide encoded by a PDE4D gene. For example, if the mutation is a frame shift mutation, the frame shift can result in a change in the encoded amino acids, and/or can result in the generation of a premature stop codon, causing generation of a truncated polypeptide. Alternatively, a polymorphism associated with a susceptibility to stroke can be a synonymous

10   mutation in one or more nucleotides (i.e., a mutation that does not result in a change in the polypeptide encoded by a PDE4D gene). Such a polymorphism may alter splicing sites, affect the stability or transport of mRNA, or otherwise affect the transcription or translation of the gene. A PDE4D gene that has any of the mutations described above is referred to herein as a "mutant gene."

15   In a first method of diagnosing a susceptibility to stroke, hybridization methods, such as Southern analysis, Northern analysis, or *in situ* hybridizations, can be used (see Current Protocols in Molecular Biology, Ausubel, F. *et al.*, eds., John Wiley & Sons, including all supplements through 1999). For example, a biological sample from a test subject (a "test sample") of genomic DNA, RNA, or cDNA, is

20   obtained from an individual suspected of having, being susceptible to or predisposed for, or carrying a defect for, stroke (the "test individual"). The individual can be an adult, child, or fetus. The test sample can be from any source which contains genomic DNA, such as a blood sample, sample of amniotic fluid, sample of cerebrospinal fluid, or tissue sample from skin, muscle, buccal or conjunctival

25   mucosa, placenta, gastrointestinal tract or other organs. A test sample of DNA from fetal cells or tissue can be obtained by appropriate methods, such as by amniocentesis or chorionic villus sampling. The DNA, RNA, or cDNA sample is then examined to determine whether a polymorphism in *PDE4D* is present, and/or to determine which splicing variant(s) encoded by PDE4D is present. The presence of

30   the polymorphism or splicing variant(s) can be indicated by hybridization of the gene in the genomic DNA, RNA, or cDNA to a nucleic acid probe. A "nucleic acid

probe", as used herein, can be a DNA probe or an RNA probe; the nucleic acid probe can contain at least one polymorphism in PDE4D or contains a nucleic acid encoding a particular splicing variant of PDE4D. The probe can be any of the nucleic acid molecules described above (e.g., the gene, a fragment, a vector

5   comprising the gene, a probe or primer, etc.).

To diagnose a susceptibility to stroke, a hybridization sample is formed by contacting the test sample containing PDE4D, with at least one nucleic acid probe. A preferred probe for detecting mRNA or genomic DNA is a labeled nucleic acid probe capable of hybridizing to mRNA or genomic DNA sequences described

10   herein. The nucleic acid probe can be, for example, a full-length nucleic acid molecule, or a portion thereof, such as an oligonucleotide of at least 15, 30, 50, 100, 250 or 500 nucleotides in length and sufficient to specifically hybridize under stringent conditions to appropriate mRNA or genomic DNA. For example, the nucleic acid probe can be all or a portion of SEQ ID NO: 1 which may optionally

15   comprise at least one polymorphism shown in Tables 9 and 10 , or the complement thereof, or a portion thereof; or can be a nucleic acid encoding a portion of SEQ ID NO: 2, 3, 4, 5, 6, 7, 8, 9, 10, 12 or 14. Other suitable probes for use in the diagnostic assays of the invention are described above (see e.g., probes and primers discussed under the heading, "Nucleic Acids of the Invention").

20   The hybridization sample is maintained under conditions which are sufficient to allow specific hybridization of the nucleic acid probe to PDE4D. "Specific hybridization", as used herein, indicates exact hybridization (e.g., with no mismatches). Specific hybridization can be performed under high stringency conditions or moderate stringency conditions, for example, as described above. In a

25   particularly preferred embodiment, the hybridization conditions for specific hybridization are high stringency.

Specific hybridization, if present, is then detected using standard methods. If specific hybridization occurs between the nucleic acid probe and PDE4D in the test sample, then PDE4D has the polymorphism, or is the splicing variant, that is present

30   in the nucleic acid probe. More than one nucleic acid probe can also be used concurrently in this method. Specific hybridization of any one of the nucleic acid

probes is indicative of a polymorphism in PDE4D, or of the presence of a particular splicing variant encoding PDE4D and is therefore diagnostic for a susceptibility to stroke.

In Northern analysis (see Current Protocols in Molecular Biology, Ausubel,

5   F. *et al.*, eds., John Wiley & Sons, *supra*) the hybridization methods described above are used to identify the presence of a polymorphism or a particular splicing variant, associated with a susceptibility to stroke. For Northern analysis, a test sample of RNA is obtained from the individual by appropriate means. Specific hybridization of a nucleic acid probe, as described above, to RNA from the individual is indicative

10   of a polymorphism in PDE4D, or of the presence of a particular splicing variant encoded by PDE4D, and is therefore diagnostic for a susceptibility to stroke.

For representative examples of use of nucleic acid probes, see, for example, U.S. Patents No. 5,288,611 and 4,851,330.

Alternatively, a peptide nucleic acid (PNA) probe can be used instead of a

15   nucleic acid probe in the hybridization methods described above. PNA is a DNA mimic having a peptide-like, inorganic backbone, such as N-(2-aminoethyl)glycine units, with an organic base (A, G, C, T or U) attached to the glycine nitrogen via a methylene carbonyl linker (see, for example, Nielsen, P.E. *et al., Bioconjugate Chemistry*, 1994, 5, American Chemical Society, p. 1 (1994). The PNA probe can

20   be designed to specifically hybridize to a gene having a polymorphism associated with a susceptibility to stroke. Hybridization of the PNA probe to PDE4D is diagnostic for a susceptibility to stroke.

In another method of the invention, mutation analysis by restriction digestion can be used to detect a mutant gene, or genes containing a polymorphism(s), if the

25   mutation or polymorphism in the gene results in the creation or elimination of a restriction site. A test sample containing genomic DNA is obtained from the individual. Polymerase chain reaction (PCR) can be used to amplify PDE4D (and, if necessary, the flanking sequences) in the test sample of genomic DNA from the test individual. RFLP analysis is conducted as described (see Current Protocols in

30   Molecular Biology, *supra*). The digestion pattern of the relevant DNA fragment

indicates the presence or absence of the mutation or polymorphism in PDE4D, and

therefore indicates the presence or absence of this susceptibility to stroke.

Sequence analysis can also be used to detect specific polymorphisms in

PDE4D. A test sample of DNA or RNA is obtained from the test individual. PCR

5   or other appropriate methods can be used to amplify the gene, and/or its flanking

sequences, if desired. The sequence of PDE4D, or a fragment of the gene, or cDNA,

or fragment of the cDNA, or mRNA, or fragment of the mRNA, is determined, using

standard methods. The sequence of the gene, gene fragment, cDNA, cDNA

fragment, mRNA, or mRNA fragment is compared with the known nucleic acid

10   sequence of the gene, cDNA (e.g., SEQ ID NO:1 which may optionally comprise at

least one polymorphism shown in Tables 9 and 10, or a nucleic acid sequence

encoding SEQ ID NO: 2, 3, 4, 5, 6, 7, 8, 9, 10, 12 or 14, or a fragment thereof) or

mRNA, as appropriate. The presence of a polymorphism in PDE4D indicates that

the individual has a susceptibility to stroke.

15   Allele-specific oligonucleotides can also be used to detect the presence of a

polymorphism in PDE4D, through the use of dot-blot hybridization of amplified

oligonucleotides with allele-specific oligonucleotide (ASO) probes (see, for

example, Saiki, R. *et al.*, (1986), *Nature (London) 324*:163-166). An "allele-specific

oligonucleotide" (also referred to herein as an "allele-specific oligonucleotide

20   probe") is an oligonucleotide of approximately 10-50 base pairs, preferably

approximately 15-30 base pairs, that specifically hybridizes to PDE4D, and that

contains a polymorphism associated with a susceptibility to stroke. An allele-

specific oligonucleotide probe that is specific for particular polymorphisms in

PDE4D can be prepared, using standard methods (see Current Protocols in

25   Molecular Biology, *supra*). To identify polymorphisms in the gene that are

associated with a susceptibility to stroke, a test sample of DNA is obtained from the

individual. PCR can be used to amplify all or a fragment of PDE4D, and its

flanking sequences. The DNA containing the amplified PDE4D (or fragment of the

gene) is dot-blotted, using standard methods (see Current Protocols in Molecular

30   Biology, *supra*), and the blot is contacted with the oligonucleotide probe. The

presence of specific hybridization of the probe to the amplified PDE4D is then

detected. Specific hybridization of an allele-specific oligonucleotide probe to DNA from the individual is indicative of a polymorphism in PDE4D, and is therefore indicative of a susceptibility to stroke.

In another embodiment, arrays of oligonucleotide probes that are

5 complementary to target nucleic acid sequence segments from an individual, can be used to identify polymorphisms in PDE4D. For example, in one embodiment, an oligonucleotide array can be used. Oligonucleotide arrays typically comprise a plurality of different oligonucleotide probes that are coupled to a surface of a substrate in different known locations. These oligonucleotide arrays, also described

10 as "Genechips.TM.," have been generally described in the art, for example, U.S. Pat. No. 5,143,854 and PCT patent publication Nos. WO 90/15070 and 92/10092. These arrays can generally be produced using mechanical synthesis methods or light directed synthesis methods which incorporate a combination of photolithographic methods and solid phase oligonucleotide synthesis methods. See Fodor *et al.*,

15 *Science*, 251:767-777 (1991), Pirrung *et al.*, U.S. Pat. No. 5,143,854 (see also PCT Application No. WO 90/15070) and Fodor *et al.*, PCT Publication No. WO 92/10092 and U.S. Pat. No. 5,424,186, the entire teachings of each of which are incorporated by reference herein. Techniques for the synthesis of these arrays using mechanical synthesis methods are described in, e.g., U.S. Pat. Nos. 5,384,261, the

20 entire teachings of which are incorporated by reference herein.

Once an oligonucleotide array is prepared, a nucleic acid of interest is hybridized with the array and scanned for polymorphisms. Hybridization and scanning are generally carried out by methods described herein and also in, e.g., Published PCT Application Nos. WO 92/10092 and WO 95/11995, and U.S. Pat.

25 No. 5,424,186, the entire teachings of which are incorporated by reference herein. In brief, a target nucleic acid sequence which includes one or more previously identified polymorphic markers is amplified by well known amplification techniques, e.g., PCR. Typically, this involves the use of primer sequences that are complementary to the two strands of the target sequence both upstream and

30 downstream from the polymorphism. Asymmetric PCR techniques may also be used. Amplified target, generally incorporating a label, is then hybridized with the

array under appropriate conditions. Upon completion of hybridization and washing of the array, the array is scanned to determine the position on the array to which the target sequence hybridizes. The hybridization data obtained from the scan is typically in the form of fluorescence intensities as a function of location on the array.

5   Although primarily described in terms of a single detection block, e.g., for detection of a single polymorphism, arrays can include multiple detection blocks, and thus be capable of analyzing multiple, specific polymorphisms. In alternate arrangements, it will generally be understood that detection blocks may be grouped within a single array or in multiple, separate arrays so that varying, optimal

10   conditions may be used during the hybridization of the target to the array. For example, it may often be desirable to provide for the detection of those polymorphisms that fall within G-C rich stretches of a genomic sequence, separately from those falling in A-T rich segments. This allows for the separate optimization of hybridization conditions for each situation.

15   Additional description of use of oligonucleotide arrays for detection of polymorphisms can be found, for example, in U.S. Patents 5,858,659 and 5,837,832, the entire teachings of which are incorporated by reference herein.

Other methods of nucleic acid analysis can be used to detect polymorphisms in PDE4D or splicing variants encoding by PDE4D. Representative methods

20   include direct manual sequencing (Church and Gilbert, (1988), *Proc. Natl. Acad. Sci. USA 81*:1991-1995; Sanger, F. *et al.* (1977) *Proc. Natl. Acad. Sci. 74*:5463-5467; Beavis *et al.* U.S. Pat. No. 5,288,644); automated fluorescent sequencing; single-stranded conformation polymorphism assays (SSCP); clamped denaturing gel electrophoresis (CDGE); denaturing gradient gel electrophoresis (DGGE) (Sheffield,

25   V.C. *et al.* (19891) *Proc. Natl. Acad. Sci. USA 86*:232-236), mobility shift analysis (Orita, M. *et al.* (1989) *Proc. Natl. Acad. Sci. USA 86*:2766-2770), restriction enzyme analysis (Flavell *et al.* (1978) *Cell 15*:25; Geever, *et al.* (1981) *Proc. Natl. Acad. Sci. USA 78*:5081); heteroduplex analysis; chemical mismatch cleavage (CMC) (Cotton *et al.* (1985) *Proc. Natl. Acad. Sci. USA 85*:4397-4401); RNase

30   protection assays (Myers, R.M. *et al.* (1985) *Science 230*:1242); use of polypeptides

which recognize nucleotide mismatches, such as *E. coli* mutS protein; allele-specific

PCR, for example.

In another embodiment of the invention, diagnosis of a susceptibility to

stroke can also be made by examining expression and/or composition of an PDE4D

5   polypeptide, by a variety of methods, including enzyme linked immunosorbent

assays (ELISAs), Western blots, immunoprecipitations and immunofluorescence. A

test sample from an individual is assessed for the presence of an alteration in the

expression and/or an alteration in composition of the polypeptide encoded by

PDE4D, or for the presence of a particular variant encoded by PDE4D. An

10   alteration in expression of a polypeptide encoded by PDE4D can be, for example, an

alteration in the quantitative polypeptide expression (i.e., the amount of polypeptide

produced); an alteration in the composition of a polypeptide encoded by PDE4D is

an alteration in the qualitative polypeptide expression (e.g., expression of a mutant

PDE4D polypeptide or of a different splicing variant). In a preferred embodiment,

15   diagnosis of a susceptibility to stroke is made by detecting a particular splicing

variant encoded by PDE4D, or a particular pattern of splicing variants.

Both such alterations (quantitative and qualitative) can also be present. An

"alteration" in the polypeptide expression or composition, as used herein, refers to an

alteration in expression or composition in a test sample, as compared with the

20   expression or composition of polypeptide by PDE4D in a control sample. A control

sample is a sample that corresponds to the test sample (e.g., is from the same type of

cells), and is from an individual who is not affected by stroke. An alteration in the

expression or composition of the polypeptide in the test sample, as compared with

the control sample, is indicative of a susceptibility to stroke. Similarly, the presence

25   of one or more different splicing variants in the test sample, or the presence of

significantly different amounts of different splicing variants in the test sample, as

compared with the control sample, is indicative of a susceptibility to stroke. Various

means of examining expression or composition of the polypeptide encoded by

PDE4D can be used, including spectroscopy, colorimetry, electrophoresis,

30   isoelectric focusing, and immunoassays (e.g., David *et al.*, U.S. Pat. No. 4,376,110)

such as immunoblotting (see also Current Protocols in Molecular Biology,

particularly chapter 10). For example, in one embodiment, an antibody capable of

binding to the polypeptide (e.g., as described above), preferably an antibody with a

detectable label, can be used. Antibodies can be polyclonal, or more preferably,

monoclonal. An intact antibody, or a fragment thereof (e.g., Fab or F(ab')$_2$) can be

5   used. The term "labeled", with regard to the probe or antibody, is intended to

encompass direct labeling of the probe or antibody by coupling (i.e., physically

linking) a detectable substance to the probe or antibody, as well as indirect labeling

of the probe or antibody by reactivity with another reagent that is directly labeled.

Examples of indirect labeling include detection of a primary antibody using a

10   fluorescently labeled secondary antibody and end-labeling of a DNA probe with

biotin such that it can be detected with fluorescently labeled streptavidin.

   Western blotting analysis, using an antibody as described above that

specifically binds to a polypeptide encoded by a mutant PDE4D, or an antibody that

specifically binds to a polypeptide encoded by a non-mutant gene, or an antibody

15   that specifically binds to a particular splicing variant encoded by PDE4D, can be

used to identify the presence in a test sample of a particular splicing variant or of a

polypeptide encoded by a polymorphic or mutant PDE4D, or the absence in a test

sample of a particular splicing variant or of a polypeptide encoded by a non-

polymorphic or non-mutant gene. The presence of a polypeptide encoded by a

20   polymorphic or mutant gene, or the absence of a polypeptide encoded by a non-

polymorphic or non-mutant gene, is diagnostic for a susceptibility to stroke, as is the

presence (or absence) of particular splicing variants encoded by the PDE4D gene.

   In one embodiment of this method, the level or amount of polypeptide

encoded by PDE4D in a test sample is compared with the level or amount of the

25   polypeptide encoded by PDE4D in a control sample. A level or amount of the

polypeptide in the test sample that is higher or lower than the level or amount of the

polypeptide in the control sample, such that the difference is statistically significant,

is indicative of an alteration in the expression of the polypeptide encoded by

PDE4D, and is diagnostic for a susceptibility to stroke. Alternatively, the

30   composition of the polypeptide encoded by PDE4D in a test sample is compared

with the composition of the polypeptide encoded by PDE4D in a control sample

(e.g., the presence of different splicing variants). A difference in the composition of the polypeptide in the test sample, as compared with the composition of the polypeptide in the control sample, is diagnostic for a susceptibility to stroke. In another embodiment, both the level or amount and the composition of the

5   polypeptide can be assessed in the test sample and in the control sample. A difference in the amount or level of the polypeptide in the test sample, compared to the control sample; a difference in composition in the test sample, compared to the control sample; or both a difference in the amount or level, and a difference in the composition, is indicative of a susceptibility to stroke.

10   Kits (e.g., reagent kits) useful in the methods of diagnosis comprise components useful in any of the methods described herein, including for example, hybridization probes or primers as decribed herein (e.g., labeled probes or prmers), reagents for detection of labeled molecules, restriction enzymes (e.g., for RFLP analysis), allele-specific oligonucleotides, antibodies which bind to mutant or to

15   non-mutant (native) PDE4D polypeptide, means for amplification of nucleic acids comprising PDE4D, or means for analyzing the nucleic acid sequence of PDE4D or for analyzing the amino acid sequence of an PDE4D polypeptide, etc.

        SCREENING ASSAYS AND AGENTS IDENTIFIED THEREBY

        The invention provides methods (also referred to herein as "screening

20   assays") for identifying the presence of a nucleotide that hybridizes to a nucleic acid of the invention, as well as for identifying the presence of a polypeptide encoded by a nucleic acid of the invention. In one embodiment, the presence (or absence) of a nucleic acid molecule of interest (e.g., a nucleic acid that has significant homology with a nucleic acid of the invention) in a sample can be assessed by contacting the

25   sample with a nucleic acid comprising a nucleic acid of the invention (e.g., a nucleic acid having the sequence of SEQ ID NO: 1 which may optionally comprise at least one polymorphism shown in Tables 9 and 10, or the complement thereof, or a nucleic acid encoding an amino acid having the sequence of SEQ ID NO: 2, 3, 4, 5, 6, 7, 8, 9, 10, 12 or 14, or a fragment or variant of such nucleic acids), under

30   stringent conditions as described above, and then assessing the sample for the

presence (or absence) of hybridization.   In a preferred embodiment, high stringency

conditions are conditions appropriate for selective hybridization. In another

embodiment, a sample containing the nucleic acid molecule of interest is contacted

with a nucleic acid containing a contiguous nucleotide sequence (e.g., a primer or a

5   probe as described above) that is at least partially complementary to a part of the

nucleic acid molecule of interest (e.g., a PDE4D nucleic acid), and the contacted

sample is assessed for the presence or absence of hybridization.   In a preferred

embodiment, the nucleic acid containing a contiguous nucleotide sequence is

completely complementary to a part of the nucleic acid molecule of interest.

10   In any of these embodiment, all or a portion of the nucleic acid of interest

can be subjected to amplification prior to performing the hybridization.

             In another embodiment, the presence (or absence) of a polypeptide of

interest, such as a polypeptide of the invention or a fragment or variant thereof, in a

sample can be assessed by contacting the sample with an antibody that specifically

15   hybridizes to the polypeptide of interest (e.g., an antibody such as those described

above), and then assessing the sample for the presence (or absence) of binding of the

antibody to the polypeptide of interest.

             In another embodiment, the invention provides methods for identifying

agents (e.g., fusion proteins, polypeptides, peptidomimetics, prodrugs, receptors,

20   binding agents, antibodies, small molecules or other drugs, or ribozymes which alter

(e.g., increase or decrease) the activity of the polypeptides described herein, or

which otherwise interact with the polypeptides herein. For example, such agents can

be agents which bind to polypeptides described herein (e.g., PDE4D binding agents);

which have a stimulatory or inhibitory effect on, for example, activity of

25   polypeptides of the invention; or which change (e.g., enhance or inhibit) the ability

of the polypeptides of the invention to interact with PDE4D binding agents (e.g.,

receptors or other binding agents); or which alter posttranslational processing of the

PDE4D polypeptide (e.g., agents that alter proteolytic processing to direct the

polypeptide from where it is normally synthesized to another location in the cell,

30   such as the cell surface; agents that alter proteolytic processing such that more

polypeptide is released from the cell, etc.

In one embodiment, the invention provides assays for screening candidate or test agents that bind to or modulate the activity of polypeptides described herein (or biologically active portion(s) thereof), as well as agents identifiable by the assays. Test agents can be obtained using any of the numerous approaches in combinatorial

5 library methods known in the art, including: biological libraries; spatially addressable parallel solid phase or solution phase libraries; synthetic library methods requiring deconvolution; the 'one-bead one-compound' library method; and synthetic library methods using affinity chromatography selection. The biological library approach is limited to polypeptide libraries, while the other four approaches are

10 applicable to polypeptide, non-peptide oligomer or small molecule libraries of compounds (Lam, K.S. (1997) *Anticancer Drug Des., 12*:145).

In one embodiment, to identify agents which alter the activity of a PDE4D polypeptide, a cell, cell lysate, or solution containing or expressing a PDE4D polypeptide (e.g., SEQ ID NO: 2, 3, 4, 5, 6, 7, 8, 9, 10, 12 or 14, or another splicing

15 variant encoded by PDE4D), or a fragment or derivative thereof (as described above), can be contacted with an agent to be tested; alternatively, the polypeptide can be contacted directly with the agent to be tested. The level (amount) of PDE4D activity is assessed (e.g., the level (amount) of PDE4D activity is measured, either directly or indirectly), and is compared with the level of activity in a control (i.e., the

20 level of activity of the PDE4D polypeptide or active fragment or derivative thereof in the absence of the agent to be tested). If the level of the activity in the presence of the agent differs, by an amount that is statistically significant, from the level of the activity in the absence of the agent, then the agent is an agent that alters the activity of PDE4D polypeptide. An increase in the level of PDE4D activity relative to a

25 control, indicates that the agent is an agent that enhances (is an agonist of) PDE4D activity. Similarly, a decrease in the level of PDE4D activity relative to a control, indicates that the agent is an agent that inhibits (is an antagonist of) PDE4D activity. In another embodiment, the level of activity of a PDE4D polypeptide or derivative or fragment thereof in the presence of the agent to be tested, is compared with a

30 control level that has previously been established. A level of the activity in the

presence of the agent that differs from the control level by an amount that is statistically significant indicates that the agent alters PDE4D activity.

The present invention also relates to an assay for identifying agents which alter the expression of the PDE4D gene (e.g., antisense nucleic acids, fusion 5 proteins, polypeptides, peptidomimetics, prodrugs, receptors, binding agents, antibodies, small molecules or other drugs, or ribozymes) which alter (e.g., increase or decrease) expression (e.g., transcription or translation) of the gene or which otherwise interact with the nucleic acids described herein, as well as agents identifiable by the assays. For example, a solution containing a nucleic acid 10 encoding PDE4D polypeptide (e.g., PDE4D gene) can be contacted with an agent to be tested. The solution can comprise, for example, cells containing the nucleic acid or cell lysate containing the nucleic acid; alternatively, the solution can be another solution which comprises elements necessary for transcription/translation of the nucleic acid. Cells not suspended in solution can also be employed, if desired. The 15 level and/or pattern of PDE4D expression (e.g., the level and/or pattern of mRNA or of protein expressed, such as the level and/or pattern of different splicing variants) is assessed, and is compared with the level and/or pattern of expression in a control (i.e., the level and/or pattern of the PDE4D expression in the absence of the agent to be tested). If the level and/or pattern in the presence of the agent differs, by an 20 amount or in a manner that is statistically significant, from the level and/or pattern in the absence of the agent, then the agent is an agent that alters the expression of PDE4D. Enhancement of PDE4D expression indicates that the agent is an agonist of PDE4D activity. Similarly, inhibition of PDE4D expression indicates that the agent is an antagonist of PDE4D activity. In another embodiment, the level and/or 25 pattern of PDE4D polypeptide(s)(e.g., different splicing variants) in the presence of the agent to be tested, is compared with a control level and/or pattern that has previously been established. A level and/or pattern in the presence of the agent that differs from the control level and/or pattern by an amount or in a manner that is statistically significant indicates that the agent alters PDE4D expression.

30   In another embodiment of the invention, agents which alter the expression of the PDE4D gene or which otherwise interact with the nucleic acids described herein,

can be identified using a cell, cell lysate, or solution containing a nucleic acid encoding the promoter region of the PDE4D gene operably linked to a reporter gene. After contact with an agent to be tested, the level of expression of the reporter gene (e.g., the level of mRNA or of protein expressed) is assessed, and is compared with

5　the level of expression in a control (i.e., the level of the expression of the reporter gene in the absence of the agent to be tested). If the level in the presence of the agent differs, by an amount or in a manner that is statistically significant, from the level in the absence of the agent, then the agent is an agent that alters the expression of PDE4D, as indicated by its ability to alter expression of a gene that is operably

10　linked to the PDE4D gene promoter. Enhancement of the expression of the reporter indicates that the agent is an agonist of PDE4D activity. Similarly, inhibition of the expression of the reporter indicates that the agent is an antagonist of PDE4D activity. In another embodiment, the level of expression of the reporter in the presence of the agent to be tested, is compared with a control level that has

15　previously been established. A level in the presence of the agent that differs from the control level by an amount or in a manner that is statistically significant indicates that the agent alters PDE4D expression.

　　　　Agents which alter the amounts of different splicing variants encoded by PDE4D (e.g., an agent which enhances activity of a first splicing variant, and which

20　inhibits activity of a second splicing variant), as well as agents which are agonists of activity of a first splicing variant and antagonists of activity of a second splicing variant, can easily be identified using these methods described above.

　　　　In other embodiments of the invention, assays can be used to assess the impact of a test agent on the activity of a polypeptide in relation to a PDE4D binding

25　agent. For example, a cell that expresses a compound that interacts with PDE4D (herein referred to as a "PDE4D binding agent", which can be a polypeptide or other molecule that interacts with PDE4D, such as a receptor) is contacted with PDE4D in the presence of a test agent, and the ability of the test agent to alter the interaction between PDE4D and the PDE4D binding agent is determined. Alternatively, a cell

30　lysate or a solution containing the PDE4D binding agent, can be used. An agent

which binds to PDE4D or the PDE4D binding agent can alter the interaction by interfering with, or enhancing the ability of PDE4D to bind to, associate with, or otherwise interact with the PDE4D binding agent. Determining the ability of the test agent to bind to PDE4D or an PDE4D binding agent can be accomplished, for

5   example, by coupling the test agent with a radioisotope or enzymatic label such that binding of the test agent to the polypeptide can be determined by detecting the labeled with ^{125}I, ^{35}S, ^{14}C or ^{3}H, either directly or indirectly, and the radioisotope detected by direct counting of radioemmission or by scintillation counting. Alternatively, test agents can be enzymatically labeled with, for example,

10   horseradish peroxidase, alkaline phosphatase, or luciferase, and the enzymatic label detected by determination of conversion of an appropriate substrate to product. It is also within the scope of this invention to determine the ability of a test agent to interact with the polypeptide without the labeling of any of the interactants. For example, a microphysiometer can be used to detect the interaction of a test agent

15   with PDE4D or a PDE4D binding agent without the labeling of either the test agent, PDE4D, or the PDE4D binding agent. McConnell, H.M. et al. (1992) Science, 257:1906-1912. As used herein, a "microphysiometer" (e.g., Cytosensor™) is an analytical instrument that measures the rate at which a cell acidifies its environment using a light-addressable potentiometric sensor (LAPS). Changes in this

20   acidification rate can be used as an indicator of the interaction between ligand and polypeptide. See the Examples Section for a discussion of know PDE4D binding partners. Thus, these receptors can be used to screen for compounds that are PDE4D receptor agonists for use in treating stroke or PDE4D receptor antagonists for studying stroke. The linkage data provided herein, for the first time, provides such

25   connection to stroke. Drugs could be designed to regulate PDE4D receptor activation which in turn can be used to regulate signaling pathways and transcription events of genes downstream, such as Cbfa1.

In another embodiment of the invention, assays can be used to identify polypeptides that interact with one or more PDE4D polypeptides, as described

30   herein. For example, a yeast two-hybrid system such as that described by Fields and Song (Fields, S. and Song, O., Nature 340:245-246 (1989)) can be used to identify

polypeptides that interact with one or more PDE4D polypeptides. ⌐ such a yeast

two-hybrid system, vectors are constructed based on the flexibility of a transcription

factor which has two functional domains (a DNA binding domain and a transcription

activation domain). If the two domains are separated but fused to two different

5   proteins that interact with one another, transcriptional activation can be achieved,

and transcription of specific markers (e.g., nutritional markers such as His and Ade,

or color markers such as lacZ) can be used to identify the presence of interaction and

transcriptional activation. For example, in the methods of the invention, a first

vector is used which includes a nucleic acid encoding a DNA binding domain and

10   also an PDE4D polypeptide, splicing variant, or fragment or derivative thereof, and

a second vector is used which includes a nucleic acid encoding a transcription

activation domain and also a nucleic acid encoding a polypeptide which potentially

may interact with the PDE4D polypeptide, splicing variant, or fragment or derivative

thereof (e.g., a PDE4D polypeptide binding agent or receptor). Incubation of yeast

15   containing the first vector and the second vector under appropriate conditions (e.g.,

mating conditions such as used in the Matchmaker™ system from Clontech) allows

identification of colonies which express the markers of interest. These colonies can

be examined to identify the polypeptide(s) which interact with the PDE4D

polypeptide or fragment or derivative thereof. Such polypeptides may be useful as

20   agents which alter the activity of expression of an PDE4D polypeptide, as described

above.

      In more than one embodiment of the above assay methods of the present

invention, it may be desirable to immobilize either PDE4D, the PDE4D binding

agent, or other components of the assay on a solid support, in order to facilitate

25   separation of complexed from uncomplexed forms of one or both of the

polypeptides, as well as to accommodate automation of the assay. Binding of a test

agent to the polypeptide, or interaction of the polypeptide with a binding agent in the

presence and absence of a test agent, can be accomplished in any vessel suitable for

containing the reactants. Examples of such vessels include microtitre plates, test

30   tubes, and micro-centrifuge tubes. In one embodiment, a fusion protein (e.g., a

glutathione-S-transferase fusion protein) can be provided which adds a domain that

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allows PDE4D or a PDE4D binding agent to be bound to a matrix or other solid support.

In another embodiment, modulators of expression of nucleic acid molecules of the invention are identified in a method wherein a cell, cell lysate, or solution

5   containing a nucleic acid encoding PDE4D is contacted with a test agent and the expression of appropriate mRNA or polypeptide (e.g., splicing variant(s)) in the cell, cell lysate, or solution, is determined. The level of expression of appropriate mRNA or polypeptide(s) in the presence of the test agent is compared to the level of expression of mRNA or polypeptide(s) in the absence of the test agent. The test

10 agent can then be identified as a modulator of expression based on this comparison. For example, when expression of mRNA or polypeptide is greater (statistically significantly greater) in the presence of the test agent than in its absence, the test agent is identified as a stimulator or enhancer of the mRNA or polypeptide expression. Alternatively, when expression of the mRNA or polypeptide is less

15 (statistically significantly less) in the presence of the test agent than in its absence, the test agent is identified as an inhibitor of the mRNA or polypeptide expression. The level of mRNA or polypeptide expression in the cells can be determined by methods described herein for detecting mRNA or polypeptide.

This invention further pertains to novel agents identified by the

20 above-described screening assays. Accordingly, it is within the scope of this invention to further use an agent identified as described herein in an appropriate animal model. For example, an agent identified as described herein (e.g., a test agent that is a modulating agent, an antisense nucleic acid molecule, a specific antibody, or a polypeptide-binding agent) can be used in an animal model to

25 determine the efficacy, toxicity, or side effects of treatment with such an agent. Alternatively, an agent identified as described herein can be used in an animal model to determine the mechanism of action of such an agent. Furthermore, this invention pertains to uses of novel agents identified by the above-described screening assays for treatments as described herein. In addition, an agent identified as described

30 herein can be used to alter activity of a polypeptide encoded by PDE4D, or to alter expression of PDE4D, by contacting the polypeptide or the gene (or contacting a cell

comprising the polypeptide or the gene) with the agent identified as described
herein.

PHARMACEUTICAL COMPOSITIONS

　　　　The present invention also pertains to pharmaceutical compositions
5　　comprising nucleic acids described herein, particularly nucleotides encoding the
polypeptides described herein; comprising polypeptides described herein (e.g., one
or more of SEQ ID NO: 2, 3, 4, 5, 6, 7, 8, 9, 10, 12 or 14); and/or comprising other
splicing variants encoded by PDE4D; and/or an agent that alters (e.g., enhances or
inhibits) PDE4D gene expression or PDE4D polypeptide activity as described
10　　herein. For instance, a polypeptide, protein (e.g., an PDE4D receptor), an agent that
alters PDE4D gene expression, or a PDE4D binding agent or binding partner,
fragment, fusion protein or prodrug thereof, or a nucleotide or nucleic acid construct
(vector) comprising a nucleotide of the present invention, or an agent that alters
PDE4D polypeptide activity, can be formulated with a physiologically acceptable
15　　carrier or excipient to prepare a pharmaceutical composition. The carrier and
composition can be sterile. The formulation should suit the mode of administration.

　　　　Suitable pharmaceutically acceptable carriers include but are not limited to
water, salt solutions (e.g., NaCl), saline, buffered saline, alcohols, glycerol, ethanol,
gum arabic, vegetable oils, benzyl alcohols, polyethylene glycols, gelatin,
20　　carbohydrates such as lactose, amylose or starch, dextrose, magnesium stearate, talc,
silicic acid, viscous paraffin, perfume oil, fatty acid esters, hydroxymethylcellulose,
polyvinyl pyrolidone, etc., as well as combinations thereof. The pharmaceutical
preparations can, if desired, be mixed with auxiliary agents, e.g., lubricants,
preservatives, stabilizers, wetting agents, emulsifiers, salts for influencing osmotic
25　　pressure, buffers, coloring, flavoring and/or aromatic substances and the like which
do not deleteriously react with the active agents.

　　　　The composition, if desired, can also contain minor amounts of wetting or
emulsifying agents, or pH buffering agents. The composition can be a liquid
solution, suspension, emulsion, tablet, pill, capsule, sustained release formulation, or
30　　powder. The composition can be formulated as a suppository, with traditional

binders and carriers such as triglycerides. Oral formulation can include standard carriers such as pharmaceutical grades of mannitol, lactose, starch, magnesium stearate, polyvinyl pyrollidone, sodium saccharine, cellulose, magnesium carbonate, etc.

5   Methods of introduction of these compositions include, but are not limited to, intradermal, intramuscular, intraperitoneal, intraocular, intravenous, subcutaneous, topical, oral and intranasal. Other suitable methods of introduction can also include gene therapy (as described below), rechargeable or biodegradable devices, particle acceleration devises ("gene guns") and slow release polymeric

10   devices. The pharmaceutical compositions of this invention can also be administered as part of a combinatorial therapy with other agents.

The composition can be formulated in accordance with the routine procedures as a pharmaceutical composition adapted for administration to human beings. For example, compositions for intravenous administration typically are

15   solutions in sterile isotonic aqueous buffer. Where necessary, the composition may also include a solubilizing agent and a local anesthetic to ease pain at the site of the injection. Generally, the ingredients are supplied either separately or mixed together in unit dosage form, for example, as a dry lyophilized powder or water free concentrate in a hermetically sealed container such as an ampule or sachette indicat-

20   ing the quantity of active agent. Where the composition is to be administered by infusion, it can be dispensed with an infusion bottle containing sterile pharmaceutical grade water, saline or dextrose/water. Where the composition is administered by injection, an ampule of sterile water for injection or saline can be provided so that the ingredients may be mixed prior to administration.

25   For topical application, nonsprayable forms, viscous to semi-solid or solid forms comprising a carrier compatible with topical application and having a dynamic viscosity preferably greater than water, can be employed. Suitable formulations include but are not limited to solutions, suspensions, emulsions, creams, ointments, powders, enemas, lotions, sols, liniments, salves, aerosols, etc.,

30   which are, if desired, sterilized or mixed with auxiliary agents, e.g., preservatives, stabilizers, wetting agents, buffers or salts for influencing osmotic pressure, etc. The

agent may be incorporated into a cosmetic formulation. For topical application, also suitable are sprayable aerosol preparations wherein the active ingredient, preferably in combination with a solid or liquid inert carrier material, is packaged in a squeeze bottle or in admixture with a pressurized volatile, normally gaseous propellant, e.g.,

5   pressurized air.

Agents described herein can be formulated as neutral or salt forms. Pharmaceutically acceptable salts include those formed with free amino groups such as those derived from hydrochloric, phosphoric, acetic, oxalic, tartaric acids, etc., and those formed with free carboxyl groups such as those derived from sodium,

10   potassium, ammonium, calcium, ferric hydroxides, isopropylamine, triethylamine, 2-ethylamino ethanol, histidine, procaine, etc.

The agents are administered in a therapeutically effective amount. The amount of agents which will be therapeutically effective in the treatment of a particular disorder or condition will depend on the nature of the disorder or

15   condition, and can be determined by standard clinical techniques. In addition, *in vitro* or *in vivo* assays may optionally be employed to help identify optimal dosage ranges. The precise dose to be employed in the formulation will also depend on the route of administration, and the seriousness of the symptoms of stroke, and should be decided according to the judgment of a practitioner and each patient's

20   circumstances. Effective doses may be extrapolated from dose-response curves derived from *in vitro* or animal model test systems.

The invention also provides a pharmaceutical pack or kit comprising one or more containers filled with one or more of the ingredients of the pharmaceutical compositions of the invention. Optionally associated with such container(s) can be a

25   notice in the form prescribed by a governmental agency regulating the manufacture, use or sale of pharmaceuticals or biological products, which notice reflects approval by the agency of manufacture, use of sale for human administration. The pack or kit can be labeled with information regarding mode of administration, sequence of drug administration (e.g., separately, sequentially or concurrently), or the like. The pack

30   or kit may also include means for reminding the patient to take the therapy. The pack or kit can be a single unit dosage of the combination therapy or it can be a

plurality of unit dosages. In particular, the agents can be separated, mixed together

in any combination, present in a single vial or tablet. Agents assembled in a blister

pack or other dispensing means is preferred. For the purpose of this invention, unit

dosage is intended to mean a dosage that is dependent on the individual

5   pharmacodynamics of each agent and administered in FDA approved dosages in

standard time courses.

METHODS OF THERAPY

The present invention also pertains to methods of treatment (prophylactic

10   and/or therapeutic) for stroke, particularly ischemic and TIA, using a PDE4D

therapeutic agent. A "PDE4D therapeutic agent" is an agent that alters (e.g.,

enhances or inhibits) PDE4D polypeptide activity and/or PDE4D gene expression,

as described herein (e.g., a PDE4D agonist or antagonist). PDE4D therapeutic

agents can alter PDE4D polypeptide activity or gene expression by a variety of

15   means, such as, for example, by providing additional PDE4D polypeptide or by

upregulating the transcription or translation of the PDE4D gene; by altering

posttranslational processing of the PDE4D polypeptide; by altering transcription of

PDE4D splicing variants; or by interfering with PDE4D polypeptide activity (e.g.,

by binding to a PDE4D polypeptide), or by downregulating the transcription or

20   translation of the PDE4D gene. Representative PDE4D therapeutic

agents include the following:

nucleic acids or fragments or derivatives thereof described herein,

particularly nucleotides encoding the polypeptides described herein and vectors

comprising such nucleic acids (e.g., a gene, cDNA, and/or mRNA, such as a nucleic

25   acid encoding a PDE4D polypeptide or active fragment or derivative thereof, or an

oligonucleotide; for example, SEQ ID NO: 1 which may optionally comprise at least

one polymorphism shown in Tables 9 and 10 or a nucleic acid encoding SEQ ID

NO: 2, 3, 4, 5, 6, 7, 8, 9, 10, 12 or 14, or fragments or derivatives thereof);

polypeptides described herein (e.g., one or more of SEQ ID NO: 2, 3, 4, 5, 6,

30   7, 8, 9, 10, 12 or 14, and/or other splicing variants encoded by PDE4D, or fragments

or derivatives thereof);

other polypeptides (e.g., PDE4D receptors); PDE4D binding agents; peptidomimetics; fusion proteins or prodrugs thereof; antibodies (e.g., an antibody to a mutant PDE4D polypeptide, or an antibody to a non-mutant PDE4D polypeptide, or an antibody to a particular splicing variant encoded by PDE4D, as described

5   above); ribozymes; other small molecules;

and other agents that alter (e.g., enhance or inhibit) PDE4D gene expression or polypeptide activity, or that regulate transcription of PDE4D splicing variants (e.g., agents that affect which splicing variants are expressed, or that affect the amount of each splicing variant that is expressed.

10   More than one PDE4D therapeutic agent can be used concurrently, if desired.

The PDE4D therapeutic agent that is a nucleic acid is used in the treatment of stroke. The term, "treatment" as used herein, refers not only to ameliorating symptoms associated with the disease, but also preventing or delaying the onset of the disease, and also lessening the severity or frequency of symptoms of the disease.

15   The therapy is designed to alter (e.g., inhibit or enhance), replace or supplement activity of a PDE4D polypeptide in an individual. For example, a PDE4D therapeutic agent can be administered in order to upregulate or increase the expression or availability of the PDE4D gene or of specific splicing variants of PDE4D, or, conversely, to downregulate or decrease the expression or availability of

20   the PDE4D gene or specific splicing variants of PDE4D. Upregulation or increasing expression or availability of a native PDE4D gene or of a particular splicing variant could interfere with or compensate for the expression or activity of a defective gene or another splicing variant; downregulation or decreasing expression or availability of a native PDE4D gene or of a particular splicing variant could minimize the

25   expression or activity of a defective gene or the particular splicing variant and thereby minimize the impact of the defective gene or the particular splicing variant.

The PDE4D therapeutic agent(s) are administered in a therapeutically effective amount (i.e., an amount that is sufficient to treat the disease, such as by ameliorating symptoms associated with the disease, preventing or delaying the onset

30   of the disease, and/or also lessening the severity or frequency of symptoms of the disease). The amount which will be therapeutically effective in the treatment of a

particular individual's disorder or condition will depend on the symptoms and severity of the disease, and can be determined by standard clinical techniques. In addition, *in vitro* or *in vivo* assays may optionally be employed to help identify optimal dosage ranges. The precise dose to be employed in the formulation will also

5   depend on the route of administration, and the seriousness of the disease or disorder, and should be decided according to the judgment of a practitioner and each patient's circumstances. Effective doses may be extrapolated from dose-response curves derived from *in vitro* or animal model test systems.

In one embodiment, a nucleic acid of the invention (e.g., a nucleic acid

10   encoding a PDE4D polypeptide, such as SEQ ID NO:1 which may optionally comprise at least one polymorphism shown in Tables 9 and 10; or another nucleic acid that encodes a PDE4D polypeptide or a splicing variant, derivative or fragment thereof, such as a nucleic acid encoding SEQ ID NO: 2, 3, 4, 5, 6, 7, 8, 9, 10, 12 or 14) can be used, either alone or in a pharmaceutical composition as described above.

15   For example, PDE4D or a cDNA encoding the PDE4D polypeptide, either by itself or included within a vector, can be introduced into cells (either *in vitro* or *in vivo*) such that the cells produce native PDE4D polypeptide. If necessary, cells that have been transformed with the gene or cDNA or a vector comprising the gene or cDNA can be introduced (or re-introduced) into an individual affected with the disease.

20   Thus, cells which, in nature, lack native PDE4D expression and activity, or have mutant PDE4D expression and activity, or have expression of a disease-associated PDE4D splicing variant, can be engineered to express PDE4D polypeptide or an active fragment of the PDE4D polypeptide (or a different variant of PDE4D polypeptide). In a preferred embodiment, nucleic acid encoding the PDE4D

25   polypeptide, or an active fragment or derivative thereof, can be introduced into an expression vector, such as a viral vector, and the vector can be introduced into appropriate cells in an animal. Other gene transfer systems, including viral and nonviral transfer systems, can be used. Alternatively, nonviral gene transfer methods, such as calcium phosphate coprecipitation, mechanical techniques (e.g.,

30   microinjection); membrane fusion-mediated transfer via liposomes; or direct DNA uptake, can also be used.

Alternatively, in another embodiment of the invention, a nucleic acid of the invention; a nucleic acid complementary to a nucleic acid of the invention; or a portion of such a nucleic acid (e.g., an oligonucleotide as described below), can be used in "antisense" therapy, in which a nucleic acid (e.g., an oligonucleotide) which

5   specifically hybridizes to the mRNA and/or genomic DNA of PDE4D is administered or generated *in situ*. The antisense nucleic acid that specifically hybridizes to the mRNA and/or DNA inhibits expression of the PDE4D polypeptide, e.g., by inhibiting translation and/or transcription. Binding of the antisense nucleic acid can be by conventional base pair complementarity, or, for example, in the case

10   of binding to DNA duplexes, through specific interaction in the major groove of the double helix.

An antisense construct of the present invention can be delivered, for example, as an expression plasmid as described above. When the plasmid is transcribed in the cell, it produces RNA which is complementary to a portion of the

15   mRNA and/or DNA which encodes PDE4D polypeptide. Alternatively, the antisense construct can be an oligonucleotide probe which is generated *ex vivo* and introduced into cells; it then inhibits expression by hybridizing with the mRNA and/or genomic DNA of PDE4D. In one embodiment, the oligonucleotide probes are modified oligonucleotides which are resistant to endogenous nucleases, e.g.

20   exonucleases and/or endonucleases, thereby rendering them stable *in vivo*. Exemplary nucleic acid molecules for use as antisense oligonucleotides are phosphoramidate, phosphothioate and methylphosphonate analogs of DNA (see also U.S. Pat. Nos. 5,176,996; 5,264,564; and 5,256,775). Additionally, general approaches to constructing oligomers useful in antisense therapy are also described,

25   for example, by Van der Krol *et al.* ((1988) *Biotechniques* 6:958-976); and Stein *et al.* ((1988) *Cancer Res* 48:2659-2668). With respect to antisense DNA, oligodeoxyribonucleotides derived from the translation initiation site, e.g. between the -10 and +10 regions of PDE4D sequence, are preferred.

To perform antisense therapy, oligonucleotides (mRNA, cDNA or DNA) are

30   designed that are complementary to mRNA encoding PDE4D. The antisense oligonucleotides bind to PDE4D mRNA transcripts and prevent translation.

Absolute complementarity, although preferred, is not required. a sequence "complementary" to a portion of an RNA, as referred to herein, indicates that a sequence has sufficient complementarity to be able to hybridize with the RNA, forming a stable duplex; in the case of double-stranded antisense nucleic acids, a

5 single strand of the duplex DNA may thus be tested, or triplex formation may be assayed. The ability to hybridize will depend on both the degree of complementarity and the length of the antisense nucleic acid, as described in detail above. Generally, the longer the hybridizing nucleic acid, the more base mismatches with an RNA it may contain and still form a stable duplex (or triplex, as the case may be). One

10 skilled in the art can ascertain a tolerable degree of mismatch by use of standard procedures.

The oligonucleotides used in antisense therapy can be DNA, RNA, or chimeric mixtures or derivatives or modified versions thereof, single-stranded or double-stranded. The oligonucleotides can be modified at the base moiety, sugar

15 moiety, or phosphate backbone, for example, to improve stability of the molecule, hybridization, etc. The oligonucleotides can include other appended groups such as peptides (e.g. for targeting host cell receptors *in vivo*), or agents facilitating transport across the cell membrane (see, e.g., Letsinger *et al.* (1989) *Proc. Natl. Acad. Sci. USA* 86:6553-6556; Lemaitre *et al.*, (1987), *Proc. Natl. Acad Sci. USA* 84:648-652;

20 PCT International Publication No. W088/09810) or the blood-brain barrier (see, e.g., PCT International Publication No. W089/10134), or hybridization-triggered cleavage agents (see, e.g., Krol *et al.* (1988) *BioTechniques* 6:958-976) or intercalating agents. (See, e.g., Zon, (1988), *Pharm. Res.* 5:539-549). To this end, the oligonucleotide may be conjugated to another molecule (e.g., a peptide,

25 hybridization triggered cross-linking agent, transport agent, hybridization-triggered cleavage agent).

The antisense molecules are delivered to cells which express PDE4D *in vivo*. A number of methods can be used for delivering antisense DNA or RNA to cells; e.g., antisense molecules can be injected directly into the tissue site, or modified

30 antisense molecules, designed to target the desired cells (e.g., antisense linked to peptides or antibodies that specifically bind receptors or antigens expressed on the

target cell surface) can be administered systematically. Alternatively, in a preferred

embodiment, a recombinant DNA construct is utilized in which the antisense

oligonucleotide is placed under the control of a strong promoter (e.g., pol III or pol

II). The use of such a construct to transfect target cells in the patient results in the

5   transcription of sufficient amounts of single stranded RNAs that will form

complementary base pairs with the endogenous PDE4D transcripts and thereby

prevent translation of the PDE4D mRNA. For example, a vector can be introduced

in vivo such that it is taken up by a cell and directs the transcription of an antisense

RNA. Such a vector can remain episomal or become chromosomally integrated, as

10   long as it can be transcribed to produce the desired antisense RNA. Such vectors

can be constructed by recombinant DNA technology methods standard in the art and

described above. For example, a plasmid, cosmid, YAC or viral vector can be used

to prepare the recombinant DNA construct which can be introduced directly into the

tissue site. Alternatively, viral vectors can be used which selectively infect the

15   desired tissue, in which case administration may be accomplished by another route

(e.g., systematically).

Endogenous PDE4D expression can also be reduced by inactivating or

"knocking out" PDE4D or its promoter using targeted homologous recombination

(e.g., see Smithies *et al.* (1985) *Nature* 317:230-234; Thomas & Capecchi (1987)

20   *Cell* 51:503-512; Thompson *et al.* (1989) *Cell* 5:313-321). For example, a mutant,

non-functional PDE4D (or a completely unrelated DNA sequence) flanked by DNA

homologous to the endogenous PDE4D (either the coding regions or regulatory

regions of PDE4D) can be used, with or without a selectable marker and/or a

negative selectable marker, to transfect cells that express PDE4D *in vivo*. Insertion

25   of the DNA construct, via targeted homologous recombination, results in

inactivation of PDE4D. The recombinant DNA constructs can be directly

administered or targeted to the required site *in vivo* using appropriate vectors, as

described above. Alternatively, expression of non-mutant PDE4D can be increased

using a similar method: targeted homologous recombination can be used to insert a

30   DNA construct comprising a non-mutant, functional PDE4D (e.g., a gene having

SEQ ID NO:1 which may optionally comprise at least one polymorphism shown in

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Tables 9 and 10), or a portion thereof, in place of a mutant PDE4D in the cell, as described above. In another embodiment, targeted homologous recombination can be used to insert a DNA construct comprising a nucleic acid that encodes a PDE4D polypeptide variant that differs from that present in the cell.

5   Alternatively, endogenous PDE4D expression can be reduced by targeting deoxyribonucleotide sequences complementary to the regulatory region of PDE4D (i.e., the PDE4D promoter and/or enhancers) to form triple helical structures that prevent transcription of PDE4D in target cells in the body. (See generally, Helene, C. (1991) *Anticancer Drug Des.*, 6(6):569-84; Helene, C., *et al.* (1992) *Ann, N.Y.*

10   *Acad. Sci.*, 660:27-36; and Maher, L. J. (1992) *Bioassays* 14(12):807-15). Likewise, the antisense constructs described herein, by antagonizing the normal biological activity of one of the PDE4D proteins, can be used in the manipulation of tissue, e.g. tissue differentiation, both *in vivo* and *for ex vivo* tissue cultures. Furthermore, the anti-sense techniques (e.g. microinjection of antisense molecules, or transfection

15   with plasmids whose transcripts are anti-sense with regard to a PDE4D mRNA or gene sequence) can be used to investigate role of PDE4D in developmental events, as well as the normal cellular function of PDE4D in adult tissue. Such techniques can be utilized in cell culture, but can also be used in the creation of transgenic animals.

20   In yet another embodiment of the invention, other PDE4D therapeutic agents as described herein can also be used in the treatment or prevention of stroke. The therapeutic agents can be delivered in a composition, as described above, or by themselves. They can be administered systemically, or can be targeted to a particular tissue. The therapeutic agents can be produced by a variety of means,

25   including chemical synthesis; recombinant production; *in vivo* production (e.g., a transgenic animal, such as U.S. Pat. No. 4,873,316 to Meade *et al.*), for example, and can be isolated using standard means such as those described herein.

A combination of any of the above methods of treatment (e.g., administration of non-mutant PDE4D polypeptide in conjunction with antisense therapy targeting

30   mutant PDE4D mRNA; administration of a first splicing variant encoded by PDE4D

in conjunction with antisense therapy targeting a second splicing encoded by
PDE4D), can also be used.

The invention will be further described by the following non-limiting
examples. The teachings of all publications cited herein are incorporated herein by
5   reference in their entirety.

EXAMPLES

EXAMPLE 1 IDENTIFICATION OF THE PDE4D GENE WITH LINKAGE TO
STROKE

Icelandic Stroke Patients and Phenotype Characterization

10   A population-based list containing 2543 Icelandic stroke patients, diagnosed
from 1993 through 1997, was derived from two major hospitals in Iceland and the
Icelandic Heart Association (the study was approved by the Icelandic Data
Protection Commission of Iceland and the National Bioethics Committee). Patients
with hemorrhagic stroke represented 6% of all patients (patients with the Icelandic
15   type of hereditary cerebral hemorrhage with amyloidosis and patients with
subarachnoid hemorrhage were excluded). Ischemic stroke accounted for 67% of
the total patients and TIAs 27%. The distribution of stroke suptypes in this study is
similar to that reported in other Caucasian populations (Mohr, J.P., *et al., Neurology*,
28:754-762 (1978); L. R. Caplan, *In Stroke, A Clinical Approach*
20   (Butterworth-Heinemann, Stoneham, MA, ed 3, (1993)).

The list of approximately 2000 living patients was run through our
computerized genealogy database. A comprehensive genealogy database that has
been established at deCODE genetics, Inc. was used to cluster the patients in
pedigrees. Each version of the computerized genealogy database is reversibly
25   encrypted by the Data Protection Commission of Iceland before arriving at the
laboratory (Gulcher, J.R., *et al., Eur. J. Hum. Genet. 8*:739 (2000)). The database
uses a patient list, with encrypted personal identifiers, as input, and recursive
algorithms to find all ancestors in the database who are related to any member on the

input list within a given number of generations back (Gulcher, J.R., and Stefansson, K., *Clin. Chem. Lab. Med.* 36:523 (1998)) covering the whole Icelandic nation. The cluster function then searches for ancestors who are common to any two or more members of the input list. One hundred and seventy-nine families with two or more

5 living patients were chosen for the study with a total of 476 patients connected within 6 meioses (6 meioses connect second cousins). Informed consent was obtained from all patients and their relatives whose DNA samples were used in the linkage scan. The mean separation between affected pairs is 4.8 meioses. Of the patients selected for the study 73% had ischemic strokes, 23% TIAs and 4%

10 hemorrhagic strokes.

In the selected families, hemorrhagic stroke patients clustered with ischemic stroke and TIA patients, and there were no families with a striking preponderance of hemorrhagic stroke or of the subtypes of ischemic stroke. Patients with ischemic stroke were reclassified according to the TOAST (Trial of Org 10172 in Acute

15 Stroke Treatment) sub-classification system for stroke (Adams, H.P., Jr., *et al.,* *Stroke, 24:*34-41 (1993)). This system includes five categories: (1) large-artery atherosclerosis, (2) cardioembolism, (3) small-artery occlusion (lacune), (4) stroke of other determined etiology and (5) stroke of undetermined etiology. The diagnoses were based on clinical features and on data from ancillary diagnostic studies.

20 Patients defined with large-artery atherosclerosis had clinical and brain imaging findings of cerebral cortical dysfunction and either significant (>70%) stenosis (this is a stricter criteria than used in TOAST where 50% stenosis is the cut-off) or occlusion of a major brain artery or branch cortical artery. Potential sources of cardiogenic embolism were excluded. The category cardioembolism included

25 patients with at least one cardiac source for an embolus and potential large-artery sources of thromobosis and embolism was eliminated. Patients with small-artery occlusion had one of the traditional clinical lacunar syndromes and no evidence of cerebral cortical dysfunction. Potential cardiac source of embolus and stenosis >70% in an ipsilateral extracranial artery was excluded. The category, acute stroke

30 of other determined etiology, included patients with rare causes of stroke and patients with two or more potential causes of stroke. If the causes of stroke could

not be determined despite extensive evaluation patients were included in the

category stroke of undetermined etiology. Fig. 1A and Fig. 1B display two

pedigrees each affected by several of the stroke subtypes, including hemorrhagic

stroke. Apparently what is inherited in stroke is the broadly defined phenotype.

5   *Genome-wide scan*

A genome-wide scan was performed using a framework map of about 1000

microsatellite markers. The DNA samples were genotyped using approximately

1000 fluorescently labelled primers. A microsatellite screening set based in part on

the ABI Linkage Marker (v2) screening set and the ABI Linkage Marker (v2)

10   intercalating set in combination with 500 custom-made markers were developed.

All markers were extensively tested for robustness, ease of scoring, and efficiency in

4X multiplex PCR reactions. In the framework marker set, the average spacing

between markers was approximately 4 cM with no gaps larger than 10 cM. Marker

positions were obtained from the Marshfield map

15   (*http://research.marshfieldclinic.org/genetics*) except for a three-marker putative .

inversion on chromosome 8 (Jonsdottir, G.M., *et al., Am. J. Hum. Genet., 67 (Suppl.*

2):332 (2000); Yu, A., *et al., Am. J. Hum. Genet.. 67 (Suppl. 2)*:10 (2000). The PCR

amplifications were set up, run and pooled on Perkin Elmer/Applied Biosystems 877

Integrated Catalyst Thermocyclers with a similar protocol for each marker. The

20   reaction volume used was 5 µl and for each PCR reaction 20 ng of genomic DNA

was amplified in the presence of 2 pmol of each primer, 0.25 U AMPLITAQ GOLD

(DNA polymerase; trademark of Roche Molecular Systems), 0.2 mM dNTPs and 2.5

mM MgCl2 (buffer was supplied by manufacturer). The PCR conditions used were

95°C for 10 minutes, then 37 cycles of 15 s at 94°C, 30s at 55°C and 1 min at 72°C.

25   The PCR products were supplemented with the internal size standard and the pools

were separated and detected on Applied Biosystems model 377 Sequencer using

v3.0 GENESCAN (peak calling software; trademark of Applied Biosystems).

Alleles were called automatically with the TRUEALLELE (computer program for

alleles identification; trademark of Cybegenetics, Inc.) program (*www.cybgen.com*),

30   and the program, DECODE-GT (computer editing program that works downstream

of the TRUEALLELE program; trademark of deCODE genetics, Inc.), was used to
fractionate according to quality and edit the called genotypes (Palsson, B., *et al.*,
Genome Res. 9:1002 (1999)). At least 180 Icelandic controls were genotyped to
derive allelic frequencies.

5   A total of 476 patients and 438 relatives were genotyped. The data was
analyzed and the statistical significance determined by applying affecteds-only
allele-sharing methods (which does not specify any particular inheritance model)
implemented in the ALLEGRO (computer program for multipoint linkage analysis;
trademark of deCODE genetics, Inc.) program which calculates lod scores based on
10   multipoint calculations. Our baseline linkage analysis uses the S_{pairs} scoring function
(Kruglyak, L., *et al.*, *Am. J. Hum. Genet., 58*:1347 (1996)), the exponential
allele-sharing model (Kong, A. and Cox, N.J., *Am. J. Hum. Genet., 61*:1179
(1997)), and a family weighting scheme which is halfway, on the log scale, between
weighting each affected pair equally and weighting each family equally. In the
15   analysis we treat all genotyped individuals who are not affected as "unknown". All
linkage analyses in this paper were performed using multipoint calculation with the
program ALLEGRO (deCODE genetics, Inc.) (Gudbjartsson, D.F., *et al.*, *Nat.
Genet. 25*:12 (2000)) .

The allele sharing lod scores for the genome scan using the framework map
20   showed three regions that achieved a lod score above 1.0. Two of these regions are
on chromosome 5q. The first peak is at approximately 69 cM with a lod score of
2.00. The second peak is at 99 cM with a lod score of 1.14. The third region is on
chromosome 14q at 55 cM with a lod score of 1.24.

The information for linkage at the 5q locus was increased by genotyping an
25   additional 45 markers over a 45 cM segment which spanned both peaks. The
information used here is defined by Nicolae (D. L. Nicolae, Thesis, University of
Chicago (1999)) and has been demonstrated to be asymptotically equivalent to a
classical measure of the fraction of missing information (Dempster, A.P., *et al.*, *J. R.
Statist. Soc. B, 39*:1 (1977)). While the lod score at the second peak dropped slightly
30   to around 1.05, the lod score at the first peak increased to 3.39. However, close
inspection of our results suggested that not only does the Marshfield genetic map

(*http://research.marshfieldclinic.org/genetics*) lack resolution (many markers
assigned the same map location), but also there may be some errors in their order.
As a result, the genetic length of the region estimated using our material was
substantially greater than what is reported. By modifying the ALLEGRO (deCODE

5   genetics, Inc.) program, we applied the EM algorithm to our data to estimate the
genetic distances between markers. We found that our estimate of the genetic length
of the region was substantially longer than that given in the Marshfield map. This
indicates a problem with marker order because, in general, incorrect marker order
leads to an increased number of apparent crossovers and increases the apparent

10   genetic length.

Physical and genetic mapping

        The marker order and inter-marker distances were improved by constructing
high density physical and genetic maps over a 20 cM region between markers
D5S474 and D5S2046. A combination of data from coincident hybridizations of

15   BAC membranes using a high density of STSs and the Fingerprinting Contig
database was used to build large contigs of BACs from the RPCI-11 library. The
order of the linkage markers was also confirmed by high-resolution genetic mapping
using the stroke families supplemented with over 112 other large nuclear families
(Fig. 3). High resolution genetic mapping was used both to anchor and place in

20   order contigs found by physical mapping as well as to obtain accurate inter-marker
distances for the correctly ordered markers. Data from 112 Icelandic nuclear
families (sibships with their parents, containing from two to seven siblings) were
analyzed together with the nuclear families available within the stroke pedigrees.
For the purpose of genetic mapping the 112 nuclear families alone provide 588

25   meioses, and the total number of meioses available for mapping was over 2000. By
comparison, the Marshfield genetic map was constructed based on 182 meioses.
The large number of meiotic events within our families provides the ability to map
markers to the resolution of 0.5 to 1.0 cM. Combining this information with the
physical map resulted in a highly reliable order of markers and inter-marker

30   distances within this 20 cM region. Linkage markers common to the genetic and

physical maps were used to anchor and place in order four of the physically mapped

contigs. By integrating the genetic and physical maps a most likely order of 30

polymorphic markers was derived (Fig. 3).

BAC contigs were generated by a method that combines coincident primer

5   hybridization with data mining. The RPCI-11 human male BAC library segments 1

& 2 (Pieter de Jong, Children's Hospital Oakland Research Institute) containing

about 200,000 clones with a 12X coverage, were gridded using a 6x6 double offset

pattern in 23 cm x 23 cm membranes with a BioGrid robot (Biorobotics Ltd.,

Cambridge, UK). Initially, hybridizations were performed with markers in the

10   region of interest according to their location in the Weizmann Institute Unified

Database (*http://bioinformatics.weizmann.ac.il/udb/*). Primer sequences were

analyzed and discarded according to their content of known repeats, *E. coli* and

vector sequences (the analysis was performed using software developed at deCODE

genetics). One hundred and fifty markers in the region (30 polymorphic markers

15   used in linkage and 120 generated from STSs) separated by an average of 130 kb

were used. The selected markers were used to generate two ^{32}P labelled probes, F

that contained the pooled forward primers and R that contained the pooled reverse

primers. Reading of positive signals was performed automatically from digitized

images of resulting autoradiograms by informatics tools developed at deCODE

20   genetics. The coincident signals in both hybridizations were selected as positive

clones. A set of overlapping clones was assembled through a combination of

hybridization and BAC fingerprint walking. Fingerprints of positive clones were

analyzed using the FPC database developed at the Sanger Center. Data from FPC

contigs prebuilt with a cutoff of 3e-12 and from sequence datamining was integrated

25   with the hybridization results. BACs in the region detected by data mining and

hybridization were re-arrayed using a Multiprobe Ilex robot (Packard, Meriden, CT).

Small membranes (8 cm x 12 cm) were gridded in 6x6 double offset pattern and

individually hybridized with the markers of interest. Positive patterns were

transferred using transparencies to an Excel file containing macros to provide BAC

30   to marker associations. A visual map was generated by combining the hybridization,

fingerprinting and sequence data. New markers were generated from BAC end

sequences to close the gap. After several rounds of hybridization positive BACs
were assembled into 7 contigs covering approximately 20 Mb. Thirty of the
polymorphic markers used in linkage were assigned to four of the contigs (Fig. 3).
Estimation of contig lengths and distance between markers assigned to them was
5   based on the FPC program.

Twenty - seven of our 30 linkage markers mapped to three contigs in the
October 2000 release from UCSC, the UC Santa Cruz (UCSC) draft assembly
(_http://genome.ucsc.edu/_). The marker order within the contigs is in agreement with
our order with the exception of two markers. Although the UCSC assemblies are
10   improving, some contigs have incorrect order, orientation, or contig assembly. We
believe that high resolution genetic mapping and perhaps focused hybridization
experiments are still necessary to confirm accuracy of sequence assemblies. In
addition, high resolution genetic mapping provides better estimates of inter-marker
genetic distances that are also important for linkage analysis (Halpern, J. and
15   Whittermore, A.S., _Hum. Hered. 49_:194 (1999); Daw, E.W., _et al., Genet._
Epidemiol. 19:366 (2000)).

Final linkage results and localization

Linkage analysis including genotypes from the higher density markers using
the deCODE marker order resulted in a lod score of 4.40 ($P = 3.9 \times 10^{-6}$) on
20   chromosome 5q12 at the marker D5S2080. The reported P value is part of the
output of the ALLEGRO (deCODE genetics, Inc.) program. It is obtained by
comparing the observed lod score to the distribution of the lod score calculated
under the null hypothesis of no linkage and the assumption that the descent
information is complete. In this case, it agrees very well with the P value that one
25   would obtain by large sample approximation. The allele sharing lod score is the log,
base 10, of an one-degree of freedom likelihood ratio. Hence, with a one-sided test,
a lod score of 4.03 corresponds to a Z score of sqrt(2*log(10)*4.03) = 4.31. Normal
approximation gives a P value of 8.2×10^{-6}. The locus has been designated as
STRK1. With the addition of these extra markers, it was possible to narrow down
30   the region to a segment less than 6 cM, from D5S1474 to D5S398, as defined by one

drop in lod. Analyses using the marker orders based on publicly available marker maps gave lower lod scores, ranging from 2.78 to 3.94.

To further investigate the contribution of this susceptibility locus to stroke, a range of parametric models were fitted to the data. However, all analyses were still

5   *affecteds only* in the sense that individuals were either classified as affecteds or having unknown disease status. A lod score of 4.08 was obtained with a dominant model where the allele frequency of the susceptibility gene was assumed to be 5% and carriers of the mutation were assumed to have seven-fold the risk of a non-carrier. By inspecting the individual families, no obvious correlation was seen

10   between families which contribute positively to the linkage results with the prevalence of hypertension, diabetes or hyperlipidemias. When the data were reanalyzed with the hemorrhagic stroke patients removed, the allele sharing lod score increased to 4.86 at D5S2080. Although this 0.46 increase in log score suggests that *STRK1* is involved primarily in ischemic stroke and TIAs, it is not

15   statistically significant based on simulations (one sided P equals 0.09). In order to assess whether such a change in lod score would be likely to occur by chance we selected 1000 random sets of 22 patients whose status we then changed to "unknown" in an analysis. The P value we present is the fraction of the 1000 simulations which produce a lod score increase at the peak locus equal to or greater

20   than that which we observed by changing the affection status of the 22 hemorrhagic stroke patients to "unknown".

Identification of Allelic Association

All microsatellite markers in the approx. 6 cM interval (Fig. 3, markers from D5S398 to D5S1474) were analysed with respect to allelic association.

Table 1. The association of a fixed allele, with the stroke patients compared with
population controls.

Marker	Location (cM)	Allele (A)	p-value	Risk ratio	Total no. of patients	Patients with A	Total no. of controls	Controls with A
AC022125-3	68.3	0	2.83e-03	1.28	749	412	504	251
D5S2000	68.5	0	3.26e-03	1.27	717	302	555	196
D5S2091	68.6	0	5.44e-04	1.30	757	342	534	198
D17-C	68.8	0	1.91e-03	1.34	721	436	469	249
D17-B	68.9	0	1.30e-03	1.26	680	556	509	387
AC008818-1	72.7	0	3.26e-03	1.42	739	379	619	259
D5S1990	73.9	20	3.68e-03	1.68	756	75	623	36

Comment:
The alleles have conventional values resulting after subtracting the CEPH data.

Identification of Microsatellite and SNP Haplotypes Within the Gene

Fig. 5 shows a schematic representation of the genetic map showing
microsatellite and SNP haplotypes in the region of the stroke gene. Seven
haplotypes are shown from the association study of Icelandic patients (804 patients).
5   The haplotypes indicated as SW-1 and SW-2 are from an association study on
Swedish stroke patients.

A total number of 804 Icelandic patients were analyzed for microsatellite
single marker and multimarker association. The number of controls used in the
analysis was 504. Each patient had 2 or more close relatives genotyped in order to
10   derive haplotypes. The haplotypes were derived using ALLEGRO based haplotype
analysis (results shown in Table 2).

Table 2
Icelandic Patient Association

Markers	Alleles	pAllelic	All Frq Aff	All Frq Ctrl	pCarrier	Carr Frq Aff	Carr Frq Ctrl	# aff	# ctrl
All patients (n=804)									
D5S2000	0	1.12E-04	0.24	0.18	5.36E-04	0.43	0.33	744	429
D5S2091	0	5.28E-04	0.26	0.21	6.10E-04	0.46	0.37	770	478
AC022125-3	0	5.96E-04	0.33	0.27	3.24E-04	0.55	0.45	774	489
D17-C	0	9.93E-04	0.36	0.29	0.007	0.6	0.52	756	395
AC008833-6	0	0.0013	0.67	0.61	0.018	0.88	0.84	781	472
AC008818-1	0	0.0014	0.29	0.24	7.13E-04	0.51	0.41	773	482
AC008829-5	2	0.0063	0.03	0.015	0.005	0.06	0.03	645	474
(1) D5S2000 D5S2091 D17-C D17-B	0000	0.0018	0.17	0.11	0.004	0.3	0.22	552	325
(2) D5S2091 D17-C D17-B	000	9.06E-04	0.19	0.13	0.001	0.34	0.25	597	380
(3) AC008829-5 AC008833-2 AC008833-3	20 14 6	0.0017	0.01	0.002	0.002	0.029	0.004	579	431
(4) AC022125-3 AC008833-6 D5S2000 D5S2091 D17-C	00000	0.00374	0.17	0.13	0.012	0.32	0.24	629	317
(5) D5S2071 AC008879-2 AC008818-1 AC008879-3	-2 0 0 0	0.0031	0.05	0.02	0.004	0.09	0.044	489	362
(6) AC008879-2 AC008818-1 AC008879-3	0 0 0	9.25E-04	0.29	0.23	5.82E-04	0.5	0.4	621	443
(part 7) D5S2107 AC008829-5 AC008833-2	4 2 0	0.0097	0.007	0	0.009	0.01	0	540	422

Swedish patients have also been genotyped and microsatellite single and multimarker association has been analyzed using the E-M algorithm. A total number of 943 Swedish patients (stroke patients and patients with carotid stenosis) and 322 Swedish controls were analyzed (results shown in Table 3).

Table 3
Swedish Patient Association

Markers	Alleles	pAllelic	All Frq Aff	All Frq Ctrl	# aff	# ctrl
Swedish patients (n=943)						
D5S2000	2	2.39E-03			912	318
(Sw 2) AC022125-3 AC008833-6 D5S2000 D5S2091	0 0 2 0	6.0E-03	0.035	0.014	717	284
(Sw-1) AC008804-2 D17-H D17-G D5S2080	-2 4 -2 10	2.8E-03	0.057	0.053	672	113
AC008804-2 D17-H D17-G	-4 0 -2	3.7E-03	0.056	0.033	700	123

SNP haplotypes within the PDE4D gene have been identified. A total of 95 SNP's typed for approximately 500 patients and 140 controls and E-M algorithm was used to analyze the genotype (results shown in Table 4). Selected SNP's found in excess in patients (based on the E-M algorithm) were typed for a subset of relatives in order to derive haplotypes for haplotype analysis (results are shown in Table 5). SNP haplotypes 1 and 2 are located upstream of D6 exon, SNP haplotype 3 is located upstream of D8 exon and stretches over it, SNP haplotype 4 stretches over LF1 exon.

Table 4
SNP genotype analysis based E-M algorithm

SNP haplotype	Position	Alleles in Haploytpe	pAllelic	All Frq Aff	All Frq Ctrl	#Aff	#Ctrl
SNP-1	1273143-1269965	122303	9.9E-03	0.32	0.25	505	155
SNP-2	1260358-1254849	10323	2.8E-02	0.33	0.26	631	131
SNP-3	1399767-1318510	2313002	8.9E-03	0.26	0.18	759	149
SNP-4	1422008-1410824	111330	3E-02	0.56	0.48	344	128

Table 5A
SNP haplotype analysis

SNP haplo-type	Position	Alleles in haplo-type	pAllelic	All Frq Aff	All Frq Ctrl	Carr Frq Aff	Carr Frq Ctrl	# Aff	# Ctrl
SNP-1	1273143-1269965	122303	4.27E-04	0.31	0.18	0.49	0.308	111	149
SNP-2	1260358-1254849	10323	0.0043	0.32	0.2	0.508	0.35	114	128

Table 5B
SNPs in the identified SNP haplotypes

Haplotype	SNP	Public name if available	Polymorpism	position	Allele
SNP-2	1	new	T/C	1254849	3
SNP-2	2	new	A/G	1257206	2
SNP-2	3	TSC0538885	T/C	1257624	3
SNP-2	4	new	A/C	1259581	0
SNP-2	5	rs244579	T/C	1260358	1
SNP1	1	rs35284	T/C	1269965	3
SNP1	2	rs35283	A/G	1270041	0
SNP1	3	rs35281	A/G	1270553	3
SNP1	4	rs35280	G/A	1272125	2
SNP1	5	new	A/G	1272910	2
SNP1	6	rs35279	G/C	1273143	1
SNP3	1	rs255652	A/G	1318510	2
SNP3	2	rs27547	G/A	1371388	0
SNP3	3	rs26695	G/A	1390407	0
SNP3	4	rs27773	C/T	1391020	3
SNP3	5	rs1471430	C/G	1391818	1
SNP3	6	rs26705	C/T	1392198	3
SNP3	7	rs26701	G/C	1399767	2
SNP4	1	rs464311	A/G	1410824	0
SNP4	2	rs1867725	T/C	1412604	3
SNP4	3	rs153966	T/C	1414091	3
SNP4	4	new	C/T	1414804	1

Table 6A and 6B show previously known microsatellite markers and novel microsatellites in sequence. Forward and reverse primers are shown.

Table 6A   Previously Known microsatellite markers in sequence

	Accession number	Forward primer	SEQ ID NO.	Reverse primer	SEQ ID NO.
D5S2107	GDB:614475	AGCCTTTGGGCCAACA	15	CAAACCAACAGGAGTAGTGTACTTTT	16
D5S468	GDB:593646	AAATGAATGGTAGATTTAACCTGAG	17	TGGGAAAATAAATACATGCG	18
D5S2000	GDB:608769	TTATACCAGGAGAGTAGACTTTTT	19	CATGCTAATTTCAAATATGAGAG	20
D5S2091	GDB:613806	GCATTTGTCATGTGCCA	21	GGTATTTCATTCACAGCCAGTC	22
D5S2500	GDB:683034	TTAAAGGAGTGATCTCCCCC	23	GTTACAGTACCTATGGTCATGCC	24
D5S2080	GDB:613188	GCACTGTGAATTTCAAATG	25	GTCAGGGGACTGGGAT	26
D5S2018	GDB:609957	CCTGTAAACAATGAAAACCCCACTGA	27	AGACTATGCTGTGTGTGTGCCTG	28
D5S2071	GDB:612756	TCTGGGTTTACAACCTTCAAA	29	TAACTGGCTTGGCCCG	30

Table 6B   Novel microsatellites in sequence:

	Forward primer	SEQ ID NO.	Reverse primer	SEQ ID NO.
DG5S382	CAGTAAATAGTTTGCTTCAGGCATT	31	CTCATACTCTGCGTGGCTTG	32
AC008829-5	AGGGCTAAGTGGATCACAGC	33	AGAGGGTCTTGCCACTGTGT	34
AC008833-2	TCTGCAAGACTCTCGGTGCT	35	TGCAGATCTCATATTTCCATGTTT	36
AC008833-3	TCTGCCCTTGTTCCTCATC	37	GTCAAGGGAGTGATGGCAGT	38
AC022125-3	AAAATGACTGCCTCCCACAA	39	GGGAAATCATACTGCCCTCA	40
AC008833-6	AAACATAGCCACCCTGTTGC	41	TCCAAAGCCCTTAGCTTAATCA	42
D17-C	GCTCCCTGGACTGTGGTAAA	43	GCCACATTGCTGTCACATTT	44
D17-B	TTTTTCAGGGCTGGGTAGAA	45	TCCAAAGGAAGTGAAATCAGTG	46
D17-D	CTAACCCATCCTCACCCAAT	47	TGTGGCATACAGGGAAGTGA	48
AC008804-1	GTGCTGGAATTTGGCTCCTA	49	CAAACATCATTTGCCTTGC	50
AC008804-2	TCCCAAACGATAGCTGTTGC	51	GAATTAGGACGGGCTGGCTCAA	52
AC008804-3	TTTGCATTCATCACTCATTCG	53	CCCGTAGCATCTGATCCAGT	54
D17-H	AGAAAGCTTCCCCTCCACTG	55	CATTCCAGCCTGAGCTACAA	56
D17-G	TGGGCTCCAATTATCCTTCC	57	TGCAGTTTGCACTCTCCTTG	58
AC027322-12	TTATCTGTTCCCCATGCTTTT	59	TGTTACATCTTGATCTATGACGTTT	60
AC027322-10	TGTATCCTGCATCCCCTGTT	61	GGAATAACCCAAAAGTAATTGTAGTGA	62
AC027322-9	TCGTTGCCAAGATGAAAATGA	63	AAACCTCCCTGATCATCTGAA	64
AC027322-8	ACAGAGGAGCAAAGGAATCA	65	TTGGCACGAATCACTCTCTG	66
AC027322-3	CCCCATTTGGATGATGGTAA	67	TGAGAAACATCTAACGTCTTTTTCAA	68
AC027322-5	GGCACAGATAACTGGGAAGC	69	CCCCCAAAAGTACTGCATAAA	70
DG5S397	ATGTTTGGCATTTGGTGAGGT	71	CACCTGTCCCTTTGGAGGTA	72
AC008879-2	TTTTAAACGTGAAAAGTACAAGTTGC	73	ACAAAGAGCACCTTTCCAGTG	74
AC008818-1	TGCTTGGTGAAGGAATAGCC	75	GAGCCTGGGTTCTCAGGAAT	76
AC008879-3	GGCAAGAAACAGTTTGGAGGA	77	GACTGCTGTTTGCTGGTTGA	78
AC020733-1	AAATGGCTATAAAGTGCTTTGAAC	79	CGGTCTCAACAACCAGAAACA	80
AC016591-2	CAGAAACACACAGAAGTCATTCAA	81	CAGACCCAATTAATGGCAAAA	82
DG5S405	TCTGTCTTCTTTGACCCATGAAT	83	CAACACAGCGGAGACCTCATC	84

Discussion of Stroke Locus Identification

Genealogy, a comprehensive population based list of broadly defined stroke patients and non-parametric allele sharing methods have been combined to successfully map a major gene for one of the most complex diseases known. There was no

5   correlation between the contribution of the families to the locus and hypertension, diabetes or hyperlipidemias and this locus does not match any known gene contributing to these risk factors. The types of stroke studied in this work do not reflect a rare or Icelandic-specific form of stroke; rather, the diversity of the stroke phenotypes in Icelanders as well as risk factors are similar to those of most other Caucasian

10   populations (Agnarsson, U., *et al., Ann. Intern. Med., 130*:987 (1999); Eliasson, J.H., *et al., Læknablaðið, 85*:517-25 (1999); Sveinbjörnsdottir, S., *et al.*, Systematic registration of patients with Stroke and TIA admitted to The National University Hospital, Reykjavik, Iceland, in 1997, XIII. Meeting of the Icelandic Association in Internal Medicine, Akureyri, Iceland (Læknabladid, 1998); Valdimarsson, E.M., *et al.*,

15   *Læknabladid 84*:921 (1998)).

The known genetic factors contributing to common stroke may do so indirectly by increasing the risk of some of its risk factors such as diabetes, hyperlipidemias, and hypertension. It is possible that there are genetic factors for stroke that do not influence susceptibility to the known risk factors, as has been suggested by epidemiologic studies

20   for myocardial infarction (Friedlander, Y., *et al., Br. Heart J., 53*:382 (1985); Shea, S., *et al., J. Am. Coll. Cardiol., 4*:793 (1984); Myers, R.H., *et al., Am. Heart J., 120*:963 (1990)). Epidemiological studies of the common forms of stroke have given conflicting results regarding the role of family history. Some studies have shown that parental history predicts the risk of stroke independently from conventional risk factors (Liao,

25   D., *et al., Stroke, 28*:1908 (1997); Jousilahti, P., *et al., Stroke, 28*:1361 (1997)) whereas others have failed to find evidence for such independent factors (Graffagnino, C., *Stroke, 25*:1599 (1994); Kiely, D.K., *et al., Stroke, 24*:1366 (1993); Lindenstrom, E., *et al., Neuroepidemiology, 12*:37 (1993).

The work described herein is the first reported genome scan searching for genes that contribute to stroke as defined as a public health problem. The data reported herein suggests that the mapped gene contributes directly to stroke without contributing indirectly through its known risk factors. This suggests that there may be other

5   biological pathways contributing to the pathogenesis of stroke.

EXAMPLE 2 IDENTIFICATION OF THE PDE4D GENE

Sequence of the Candidate Region

We have sequenced approximately 3 Mb of the area defined by one drop in lod (Fig. 3, the genetic map of the region). The BAC (bacterial artificial clones) sequenced

10   in house are shown in Table 7A. We also used for the assembly the following publicly available BAC sequences from GenBank listed in Table 7B for the assembly. The BAC clones we sequenced are from the RCPI-11 Human BAC library (Pieter deJong, Roswell Park). The vector used was pBACe3.6. The clones were picked into a 94 well microtiter plate containing LB/chloramphenicol (25 µg/ml)/glycerol (7.5%) and stored

15   at -80°C after a single colony has been positively identified through sequencing. The clones can then be streaked out on a LB agar plate with the appropriate antibiotic, chloramphenicol (25 µg/ml)/sucrose (5%).

Table 7A

Sequenced at Decode (BAC name)	Comment	Accession number
RP11-621C19	1	AC020733
RP11-113C1	2	
RP11-412M9	2	
RP11-151G2	2	
RP11-151F7	2	
RP11-281M3	2	
RP11-421L6	2	
RP11-68E13	2	
RP11-379P8	2	
RP11-1A7	1	AC008111
RP11-422K3	2	
RP11-116A3	2	

Key to "Comment" column:

1= This BAC has a publicly available sequence,

    it was sequenced at Decode to make sure the sequence was correct

2= Only BAC end-sequence available for this BAC publicly.

Table 7B

Sequences available from

GenBank (BAC name)	Accession number	Status of sequence
RP11-621C19	AC020733	17 unordered pieces
CTD-2003D5	AC016591	complete sequence
CTD-2210C1	AC008879	7 unordered pieces
CTD-2124H11	AC008818	complete sequence
CTD-2301A11	AC008934	complete sequence
RP11-16B11	AC011929	7 unordered pieces
CTC-261E10	AC026693	complete sequence
CTD-2027G10	AC027322	complete sequence
RP11-1A7	AC008111	8 unordered pieces
CTD-2122K7	AC012315	complete sequence
CTD-2085F10	AC008804	complete sequence
CTD-2040J22	AC008791	complete sequence
RP11-235N16	AC020975	16 ordered pieces
CTD-2146O16	AC008833	complete sequence
CTD-2084I4	AC022125	17 ordered pieces
CTD-2140K22	AC008829	26 ordered pieces
CTD-2124D11	AC020924	7 ordered pieces
RP11-731H6	AC026095	21 unordered pieces

Gene identification

The gene, human cAMP specific phosphodiesterase 4D (HPDE4D) was identified in the sequenced region (Fig. 3). Twenty-three exons have been identified,

eighteen of those have previously been published. See top of Fig. 4. Five new spliced exons have been identified (referred to as 4D6, 4D7-1, 4D7-2, 4D7-3 and 4D8) in three new isoforms (PDE4D6, PDE4D7 and PDE4D8). The genomic sequence is approximately 1,691,140 bases in length.

The exon locations are indicated in Table 8 below.

Table 8

Exon	Start	End
(New) 4D7-1	142207	142328
(New) 4D7-2	444645	444775
(New) 4D7-3	641649	641878
4D4	736254	737226
4D5	861791	862202
4D3	1044051	1044190
(New) 4D6	1273404	1273709
(New) 4D8	1354347	1355128
LF1	1414511	1414702
LF2	1436943	1436979
LF3	1472965	1473235
LF4	1449835	1449542
N3	1539259	1539302
4D1/D2	1591172	1591425
ex3	1636944	1637037
ex4	1638406	1638578
ex5	1639508	1639606
ex6	1640491	1640655
ex7	1641818	1641917
ex8	1653070	1653224
ex9	1653943	1654065
ex10	1654576	1654758
ex11	1655335	1655747

The markers showing the highest association are located within the PDE4D
(Table 1, Fig. 3 and Table 5), as follows:

AC022125-3, 21 000 bp upstream of the LF1 exon

D5S2000, 37 000 bp downstream of PDE4D6 exon

D5S2091, 30 000 bp downstream of PDE4D6 exon

D17-C, 21 000 bp upstream of PDE4D6 exon

D17-B, 31 000 bp upstream of PDE4D6 exon

AC008833-6, 35 000 bp downstream of PDE4D8 exon

AC008818-1, 3000 pb upstream of PDE4D7-1 exon

AC008829-5, 89 000 bp upstream of PDE4D1/D2 exon

Haplotype (1) and (2) are located upstream of and stretch over the PDE4D6 exon

Haplotype (3) is located upstream of and stretches over the LF2-LF4 exons

Haplotype (4) stretches over PDE4D6 and PDE4D8 exons

Haplotype (5) stretches over PDE4D7-1 to PDE4D7-3 exons

Haplotype (6) stretches over PDE4D7-1 exon

Haplotype (7) stretches over LF2-exons 11

A contig for the incomplete genomic sequence of the PDE4D gene was submitted in November 2000 (GenBank entry NT_023193 by International Human Genome Project collaborators). The size of the contig is 614 481 bp (including gaps) whereas our genomic sequence for the whole PDE4D region (i.e., from the first exon for PDE4D variant) is close to 1,700,000 bp. The contig NT_023193 comprises only 11 exons of the PDE4D gene (in Fig. 4, exons 4D1/D2 - 11) and the 5' differently spliced exons are missing in the contig (in Fig. 4, exons 4D4, 4D5, 4D3, 4D6, 4D8, 4D7-1, 4D7-2, 4D7-3, LF1, LF2, LF3 and LF4).

SNPs (single nucleotide polymorphisms) detected in the sequence and mutation analysis

Publically available and novel SNP's in the PDE4D2 gene from mutation screening of all exons are illustrated in Tables 9 and 10.

Gene Identification

The identified gene PDE4D is a member of the cyclic nucleotide phosphodiesterases (PDEs). Intracellular levels of cyclic AMP and cyclic GMP are mediated by the PDEs. Cyclic nucleotides are important second messengers that regulate and mediate a number of cellular responses to extracellular signals, such as hormones, light and neurotransmitters. Intracellular levels of cAMP play a key role in the function of inflammatory and immune cells. One of the mechanisms that mediate relaxation of vascular muscle in cerebral circulation is the production of cAMP.

PDE4D Structure and Splice Forms

Phosphodiesterases are the mammalian homolog of the "dunce" gene in *Drosophila melanogaster*, implicated in learning and memory (Davis, R.L. and B. Dauwalder, *Trends Genet., 7(7)*:224-229 (1991)). PDEs are members of a large superfamily of isoenzymes subdivided into 9 and possibily 10 distinct families (Conti, M. and S.L. Jin, *Prog. Nucleic Acid Res. Mol. Biol., 63*:1-38 (1999)), with several genes in each family and more than one isoform for each gene. The significance of the diversity of PDEs is not known but many of the isoforms differ in their biochemical properties, phosphorylation, intracellular targeting, protein-protein interactions and patterns of expression in tissues, which suggests that each of the various isoforms might have distinct functions (Bolger, G.B., *Cell Signal, 6(8)*:851-859 (1994); Conti, M., *et al., Endocr. Rev., 16(3)*:370-378 (1995)).

There are four genes that encode the type 5 PDEs (PDE4A, PDE4B, PDE4C and PDE4D), which is a group of enzymes characterized by high affinity for cAMP. The gene for PDE4D was assigned to human chromosome 5q12 (Milatovich, A., *et al., Somat. Cell Mol. Genet., 20(2)*:75-86 (1994); Szpirer, C., *et al., Cytogenet. Cell Genet., 69(1-2)*:22-14 (1995)) and 5 distinct splice variants have been characterized (the short forms PDE4D1, PDE4D2 and the long forms PDE4D3, PDE4D4, and PDE4D5) (Bolger, G.B., *et al., Biochem. J., 328(Pt.2)*:539-548 (1997)) (Fig. 4). The sequence of

the human PDE4D variants show a high degree of homology to the PDE4Ds expressed in mouse and rat. The pattern of splicing and different promoter usage is highly conserved during evolution indicating an important physiological role (Nemoz, G., et al., *FEBS Lett., 384(1)*:97-102 (1996)). The PDE4D variants are generated at two major boundaries present in the gene. The first boundary corresponds to the junction of exon 2. Differential splicing in this region generates the 2 short variants PDE4D1 (586 a.a.) and PDE4D2 (508 a.a.)(Fig. 4). This splicing boundary is conserved in mouse, rat and between different human PDE4 genes. The splicing variant PDE4D2 is generated by the removal of 256 bp from the PDE4D1 sequence. The initiation codon in the PDE4D2 variant lies within exon D1/D2. Data demonstrates that the expression of the short PDE4D variants is under the control of an internal promoter regulated by cAMP (Vicini, E. and M. Conti, *Mol. Endocrinol., 11(7)*:839-850 (1997)). The second major splicing boundary is also conserved during evolution and is identical to that described in the *Drosophila* dunce gene. Splicing occurs at the intron/exon boundary at the LF1 exon (Fig. 4).

PDE function

The PDEs serve at least four major functions in the cell. They can (1) act as effector of signal transduction by interacting with receptors and G-proteins; (2) integrate the cyclic nucleotide-dependent pathway with other signal transduction pathways; (3) function as homeostatic regulators, playing a role in feedback mechanisms controlling cyclic nucleotide levels during hormone and neurotransmitter stimulation; (4) play an important role in controlling the diffusion of cyclic nucleotides and in creating subcellular domains or channeling cyclic nucleotide signaling (Conti, M. and S.L. Jin, *Prog. Nucleic Acid Res. Mol Biol., 63*:1-38.(1999)). Inhibition of PDE has long been recognized as an effective pharmacological strategy to alter intracellular cyclic nucleotide levels (Flamm, E.S., *et al., Arch. Neurol., 32(8)*:569-71 (1975)).

It has been reported that PDE4 is the predominant isozyme regulating vascular tone mediated by cAMP hydrolysis in cerebral vessels (Willette, R.N., *et al., J. Cereb. Blood Flow Metab., 17(2)*:210-9 (1997)).

A recent study on mice with targeted disruption of PDE4D gene (Hansen, G., *et al., Proc. Natl. Acad. Sci. U S A, 97(12)*:6751-6 (2000)) has demonstrated a crucial role of PDE4D in the control of smooth muscle contraction and muscarinic cholinergic receptor signaling but not in the control of airway inflammation. The lung phenotype of the PDE4D-/- mice demonstrates that this gene plays a nonredundant role in cAMP homeostasis. There is a significant reduction in PDE activity and an increase in resting and stimulated cAMP levels in the lung, indicating that other PDE4s (or other PDEs) are not up-regulated and cannot compensate for the loss of PDE4D. These findings support that PDE4D serves a unique, nonoverlapping functions in cell signalling.

No clear link between an established inherited disorder and known PDE loci has emerged, with the exception of PDE6. Inhibitors of PDEs have been shown to affect airway responsiveness and pulmonary allergic inflammation (Schudt, C., *et al., Pulm. Pharmacol. Ther., 12(2)*:123-9 (1999)). There are reports suggesting that altered PDE4 function may be linked to nephrogenic diabetes insipidus (Takeda, S., *et al., Endocrinology, 129(1)*:287-94 (1991)) or atopic dermatitis (Chan, S.C., *et al., J. Allergy Clin. Immunol., 91(6)*:1179-88 (1993)), however no mutations have been identified. It has also been reported that that vasorelaxation modulated by PDE4 (not mentioned whether it is A, B, C or D gene family) is compromised in chronic cerebral vasospasm associated with subarachnoid hemorrhage (Willette, R.N., *et al., J. Cereb. Blood Flow Metab., 17(2)*:210-9 (1997)). PDE4D itself has not been linked to stroke before.

PDE4D expression and cellular localization

PDE4Ds are expressed in human peripheral mononuclear cells (Nemoz, G., *et al., FEBS Lett, 384(1)*:97-102 (1996)), brain (Bolger, G., *et al., Mol. Cell Biol., 13(10)*:6558-71 (1993)), heart (Kostic, M.M., *et al., J. Mol. Cell Cardiol.,*

29(11):3135-46 (1997)) and vascular smooth muscle cells (Liu, H. and D.H. Maurice, *J. Biol. Chem., 274(15)*:10557-65 (1999)).

Immunoblotting of rat brain has shown that the PDE4D3, PDE4D4 and PDE4D5 proteins are present in brain (Bolger, G.B., *et al., Biochem. J., 328(Pt 2)*:539-48 (1997)) and are expressed in cortex and cerebellum from rat (Iona, S., *et al., Mol. Pharmacol., 53(1)*:23-32 (1998)). These proteins were recovered mostly or exclusively in the particulate fraction suggesting that these forms may be targeted to insoluble cellular structures. In addition a 68 kDa protein was detected which could represent PDE4D1, PDE4D2 or both. To verify this RT-PCR was performed on mRNA from rat brain and the results showed that transcripts for PDE4D1 and 2 were present. Their data also suggests that the N-terminal regions of the PDE4D3-5, derived from alternatively spliced regions of their mRNAs, are important in determining their subcellular localization activity and differential sensitivity to inhibitors and there are indications that there is a propensity for the long PDE4D isoforms to interact with particulate fraction of the cell.

Newly identified isoforms

Five new exons have been identified. Exon D6 was identified by deCODE (in silico) and verified by RT-PCR. The four other new exons have been identified using CAP-RACE amplification from cultured cells with an "long-form 1"-specific reverse primer. Three of these exons are spliced to one another and together onto LF1 and this new isoform was given the name D7. The fourth new 5' exon was spliced by itself onto LF1 and given the name D8. These constitute two previously unknown isoforms.

In terms of genomic structure, the D7 exons extend the known 5' end of PDE4D over 590,000 bp and the D8 exon lies between two previously recognized exons. The D7 isoform has an open reading frame extending into LF1, resulting in an additional 90 amino acids at the N-terminus of the predicted protein. The D8 5' exon contains a long 5' UTR, followed by an ATG near the end of the exon that extends an ORF into LF1 and results in a novel 21 N-terminal amino acids in the predicted protein.

Table 11: New Isoforms

Isoform Name	Exon		Size	Cell line
PDE4D6	D6			
PDE4D7	D7-1	5'	122 bp	SKNAS
PDE4D7	D7-2	Internal	131bp	SKNAS
PDE4D7	D7-3	Internal	230 bp	SKNAS
PDE4D8	D8	5'	782 bp	HeLa

The sequences are as follows:

D7-1:

ATAGTTGGCGTACCCTGAGGCCTGCCAGTTCCTGCCTTAATGCATATGTAGT
CGTAATTGAGTTCTGACACGGCCTTGGATGTTTCTGTCCTAAATAGCTGACA
TTGCATCTTCAAGACTGT

D7-2:

CATTCCAGTTGGCTTTTGAGTGGATACGTGCAGTGAGATCATTGACACTGGA
AACACTAGTTCCCATTTTAATTACTTAAAACACCACGATGAAAAGAAATACC
TGTGATTTGCTTTCTCGGAGCAAAAGT

D7-3:

GCCTCTGAGGAAACACTACATTCCAGTAATGAAGAGGAAGACCCTTTCCGC
GGAATGGAACCCTATCTTGTCCGGAGACTTTCATGTCGCAATATTCAGCTTC
CCCCTCTCGCCTTCAGACAGTTGGAACAAGCTGACTTGAAAAGTGAATCAGA
GAACATTCAACGACCAACCAGCCTCCCCCTGAAGATTCTGCCGCTGATTGCT
ATCACTTCTGCAGAATCCAGTGG (SEQ. ID NO.: 11; includes D7-1, D7-2 and D7-3)

New predicted amino-terminal protein sequence from above (PDE4D7):

MKRNTCDLLSRSKSASEETLHSSNEEEDPFRGMEPYLVRRLSCRNIQLPPLAFRQ
LEQADLKSESENIQRPTSLPLKILPLIAITSAESS (90 amino acids) (SEQ ID NO.:12)

D8:

TTCTCACTGCCCTGCGGTGTTTTGAACTGCCTTCTTACAGACGTCATACAGCC
CTTGAGGAATAGTTTCTGCCTGGTGAGATTGAATGATAGTTCTCATTCACAA
AACCCTGGATTCTAAGCAGGGACACACAGAAATTACTTTCGCAGGTAAATC
AGCCCACCCAGCCAAAGTGTGGAGAGATTTGTTCCTTGGCTGACTTCTTTGC
TCCACGGAGAGGAGTGTTTTCCTGTGCTTGCCCTGAAATGGAACTTCCTTGA
CAGCTCTCCCGTGTTACAGTACCTCCCGGTCATTTTCTTTTTCTCTCTCTAC
CTGCGCTCTTCGAGTGTCAGAAACCTTTAAAGCTGTTACTATGGAATTGCAA
AAAAGAGATCAAGTGACTCTTTCACTATGCTGGTTTCCCTTGTGACCCAGAT
GAAGAATCAATTCAGAATTCAGTTCCTCCCTTGGCATTGCAAGACACAGAAG
AAACTGTCACTTCCTAACAGCCTAGTACTGGAGTAAATTCAGTATGAAGGAA
GAAAGCGCTCCTGCGTGTTAGAACCTTGCCCATGAGCTGGACCGAGGACAG
GAGATGGACTCCAGGAAAATTGGATTTCTTCAAGCAGCCTCCCTTGGAAATG
GAATATCTTTAAAATCTTCTTTGCAGAAAGACAGTTAGAATGTATTAATCAG
AATAGTTGAAGACTTATTTTCCTTTTTATTTTTTTTCAAAATGAGCATTATTAT
GAAGCCAAGATCCCGATCTACAAGTTCCCTAAGGACTGCAGAGGCAGTTTG
(SEQ ID NO.:13)

New predicted amino-terminal protein sequence from above (PDE4D8):

MSIIMKPRSRSTSSLRTAEAV (21 amino acids) (SEQ ID NO.: 14).

Expression analysis

The tissues below were examined by RT-PCR, cloning and sequencing. The presence (Pos.) or absence (-) of the isoforms transcripts is shown in tables below.

Table 12A   Original Cell Lines (SKNAS and HeLa)

	D7	D8
HeLa	-	Pos.
SkNAs	Pos.	Pos.

Table 12B   Human tissue DNA panels

cDNA panels	D7	D8
Spleen	-	Pos.
Lymph node	Pos.	Pos.
Thymus	Pos.	Pos.
Tonsil	Pos.	Pos.
Leukocytes	Pos.	Pos.
Bone marrow	Pos.	Pos.
Heart	-	Pos.
Brain	-	Pos.
Placenta	Pos.	Pos.
Lung	Pos.	Pos.
Liver	-	Pos.
Skel. muscle	-	Pos.
Kidney	Pos.	Pos.
Pancreas	-	Pos.

Table 12C   Human blood cell fractions

	D7	D8
Spleen	Pos.	Pos.
Lymph node	Pos.	Pos.
Thymus	Pos.	Pos.
Tonsil	Pos.	Pos.
Leukocytes	Pos.	-
Bone marrow	Pos.	Pos.
Fetal liver	Pos.	Pos.
Mononucl. cells resting	Pos.	Pos.
CD4Pos. resting	-	Pos.
CD8Pos. resting	-	-
CD14Pos. resting	Pos.	Pos.
CD19Pos. resting	Pos.	Pos.
Mononucl. cells activated	-	-
CD4Pos. activated	-	-
CD8Pos. activated	-	-
CD19Pos. activated	-	Pos.

Table 12D   Cultured in-house endothelial and smooth muscle cells from patients

Cell type	D1	D2	D3	D5	D6	D7	D8
Normal aorta smooth musc.	Pos.	Pos.	Pos.	Pos.	Pos.	-	-
Diseased aorta smooth musc.	Pos.	Pos.	-	Pos.	Pos.	-	Pos.
Diseased aorta smooth musc.	Pos.	Pos.	-	Pos.	Pos.	-	-
Diseased femoral smooth musc.	Pos.	Pos.	-	Pos.	Pos.	-	Pos.
Normal aortic endothelial cells	Pos.	Pos.	Pos.	Pos.	Pos.	Pos.	Pos.
Diseased aortic endothelial cells	Pos.	Pos.	-	Pos.	Pos.	-	-
Diseased femoral endothelial cells	Pos.	Pos.	-	Pos.	Pos.	-/?	-/?

Isoform specific primers were designed in order to better determine the expression of different PDE4D isoforms using RT-PCR on Epstein Barr Virus (EBV) transformed B cell lines from stroke patients and controls. The results are outlined in Tables 13A and 13B below. There is a significant difference between the expression of D3 and D7 in patients compared to controls.

Table 13A   RT-PCR on EBV transformed B stroke patient cells

Patient Cells	PDE4D*	D3	D4	D5	D6	D7	D8
P-1	Pos.	Pos.	-	Pos.	-	Pos.	Pos.
P-2	Pos.	Pos.	-	Pos.	-	Pos.	-
P-3	Pos.	-	-	Pos.	-	-	-
P-4	Pos.	Pos.	-	Pos.	-	Pos.	-
P-5	Pos.	Pos.	Pos.	Pos.	-	Pos.	-
P-6	Pos.	-	Pos.	Pos.	-	Pos.	-
P-7	Pos.	Pos.	-	Pos.	-	Pos.	-
P-8	Pos.	-	-	-	-	Pos.	-
P-9	Pos.	-	-	Pos.	-	Pos.	-
P-10	Pos.	-	-	Pos.	Pos.	Pos.	-
P-11	Pos.	-	-	Pos.	-	Pos.	-
P-12	Pos.	-	-	Pos.	-	Pos.	-
P-13	Pos.	-	-	Pos.	-	Pos.	-
P-14	Pos.	-	-	Pos.	-	Pos.	-
% expr.	100	35,7	14,3	92,8	7,1	92,8	7,1

*Primers designed for the common region of PDE4D identical for all isoforms

Table 13B   RT-PCR on EBV transformed B control cells

Control Cells	PDE4D *	D3	D4	D5	D6	D7	D8
C-1	Pos.	-	-	Pos.	-	-	Pos.
C-2	Pos.	-	-	Pos.	-	-	-
C-3	Pos.	-	-	Pos.	-	-	-
C-4	Pos.	-	-	Pos.	-	-	-
C-5	Pos.	-	-	-	-	Pos.	-
C-6	Pos.	-	-	-	-	-	-
C-7	-	-	-	Pos.	-	-	Pos.
C-8	Pos.	-	-	-	-	Pos.	-
C-8	Pos.	Pos.	-	Pos.	-	Pos.	-
C-9	Pos.	-	-	-	-	Pos.	-
C-10	Pos.	-	-	Pos.	-	Pos.	-
C-11	Pos.	-	-	Pos.	-	Pos.	-
C-12	Pos.	-	-	Pos.	-	-	-
% expr.	92,3	7,7[a]	0	69,2	0	46,2[b]	15,4

[a] $p < 0.09$ using Fisher's Exact Test.

[b] $p = 0.01$ using Fisher's Exact Test

*Primers designed for the common region of PDE4D identical for all isoforms

Table 9

Publically Available SNPS; SNP ID No. from NCBI Database

rs286155	rs40512	rs251726	rs2042315	rs1544791	rs1355099
rs286156	rs35386	rs1862589	rs918590	rs851284	rs1396473
rs2061250	rs35387	rs702556	rs918591	rs1396476	rs1369285
rs286150	rs27221	rs702554	rs918592	rs1508860	rs1435071
rs206789	rs27653	rs441391	rs1115372	rs1974850	rs1435070
rs1823062	rs26955	rs446883	rs1345782	rs2136203	rs1435083
rs1823063	rs26956	rs789615	rs1363862	rs2174994	rs991551
rs1445852	rs153031	rs401207	rs1423248	rs1508863	rs1154790
rs766119	rs185190	rs364917	rs1423246	rs1508859	rs1154789
rs956721	rs37762	rs404202	rs1862614	rs1508864	rs714291
rs248910	rs37761	rs440607	rs2194256	rs1396474	rs981760
rs248912	rs1423471	rs411255	rs889305	rs1543951	rs1369288
rs187481	rs27224	rs615429	rs2113071	rs2016324	rs977418
rs153152	rs1645013	rs789396	rs2113072	rs1995780	rs977417
rs27960	rs1423472	rs37684	rs966220	rs1508865	rs977416
rs27564	rs27220	rs1445893	rs966221	rs952110	rs1529843
rs27565	rs1423473	rs37685	rs719702	rs1533019	rs1529842
rs26948	rs149079	rs1086121	rs2113073	rs2117552	rs1435077
rs40131	rs149324	rs42222	rs2113074	rs1545069	rs1369287
rs26949	rs153067	rs37707	rs2113075	rs1545070	rs1017410
rs26950	rs40354	rs37708	rs1035512	rs973700	rs1017409
rs26954	rs26951	rs37709	rs1559277	rs1583434	rs1435076
rs26953	rs153029	rs789389	rs1981848	rs1347401	rs1435075
rs152324	rs27223	rs1423247	rs1544788	rs1949017	rs1435074
rs35385	rs27222	rs874768	rs1544790	rs723962	rs978455

rs1827340	rs159621	rs1504982	rs298084	rs298027	rs295972
rs1393083	rs159625	rs877745	rs298083	rs298028	rs295971
rs988364	rs1435072	rs877744	rs298073	rs298029	rs295970
rs1017408	rs173945	rs2164661	rs298072	rs298030	rs295969
rs2053155	rs256356	rs981230	rs298071	rs169868	rs295968
rs181923	rs185351	rs1437124	rs1421400	rs177077	rs295966
rs1546364	rs256355	rs746477	rs402874	rs298032	rs726652
rs173942	rs2067024	rs893191	rs434368	rs298033	rs295965
rs159616	rs256354	rs1992112	rs371011	rs298034	rs1307218
rs159620	rs173944	rs298102	rs298063	rs298035	rs1307217
rs1501641	rs256353	rs298101	rs298062	rs298042	rs893190
rs159619	rs986400	rs2164660	rs298061	rs298044	rs1111495
rs159614	rs1504981	rs298100	rs298060	rs298045	rs295961
rs159613	rs1120533	rs298098	rs298057	rs298046	rs295960
rs159612	rs256351	rs298096	rs298056	rs298048	rs295959
rs159611	rs190458	rs298095	rs1370230	rs298049	rs295958
rs194368	rs256352	rs298094	rs297975	rs298050	rs296410
rs661576	rs171745	rs298093	rs297974	rs298051	rs295957
rs299627	rs1157709	rs1362942	rs379578	rs298052	rs295956
rs159608	rs1910790	rs1362941	rs920190	rs298053	rs295955
rs159609	rs1910789	rs298091	rs1865962	rs190936	rs295954
rs159624	rs1504985	rs298090	rs298018	rs298017	rs295949
rs1159470	rs1008709	rs298089	rs298021	rs298016	rs295980
rs159622	rs1027747	rs298088	rs298022	rs298015	rs295979
rs256349	rs869685	rs298087	rs298023	rs298014	rs295978
rs256348	rs869686	rs1421401	rs298024	rs2053229	rs1154587
rs1501640	rs924880	rs298086	rs298025	rs295974	rs296406
rs600611	rs1504983	rs298085	rs298026	rs295973	rs296405

rs295948	rs294478	rs3757.5	rs1457111	rs171800	rs403695
rs295947	rs953302	rs37576	rs1824154	rs187716	rs403672
rs295946	rs294479	rs1876209	rs2112911	rs258110	rs372309
rs295945	rs697075	rs190486	rs1551564	rs258109	rs424839
rs295944	rs294481	rs447261	rs2034895	rs258108	rs370891
rs1395334	rs294482	rs1506558	rs2081092	rs258107	rs434183
rs295943	rs294483	rs1108916	rs2112910	rs665836	rs444552
rs1035321	rs702545	rs921942	rs918583	rs392901	rs433565
rs294494	rs294484	rs924998	rs1840838	rs383444	rs1445918
rs722923	rs294485	rs176705	rs1350298	rs662643	rs441817
rs294495	rs294486	rs1156029	rs1990985	rs670169	rs433161
rs294496	rs702544	rs1156028	rs1379297	rs525099	rs428059
rs294497	rs702543	rs931857	rs1817248	rs669240	rs434422
rs294498	rs159194	rs931856	rs244569	rs381755	rs427433
rs294499	rs40215	rs931855	rs244568	rs454702	rs391377
rs294500	rs291118	rs1506557	rs244567	rs443191	rs414746
rs294501	rs1506560	rs462930	rs244565	rs380118	rs187368
rs294503	rs37569	rs458953	rs185417	rs2168649	rs244593
rs295936	rs291119	rs174039	rs258128	rs371775	rs244592
rs1395336	rs37571	rs2174624	rs258127	rs378970	rs244591
rs1395337	rs1870077	rs2135480	rs258125	rs401013	rs244590
rs294492	rs159195	rs992726	rs1348710	rs427748	rs181736
rs159196	rs37572	rs294474	rs1348709	rs427740	rs193447
rs159197	rs37573	rs294475	rs1971061	rs378869	rs2028842
rs172362	rs167161	rs988827	rs1541673	rs1902609	rs2028841
rs37579	rs37574	rs988828	rs1541672	rs389324	rs1823068
rs721784	rs1506562	rs1350297	rs258112	rs387647	rs1823067
rs697076	rs291122	rs1457110	rs258111	rs377451	rs1823066

rs244588	rs35275	rs2014012	rs531105	rs27691	rs464311
rs168641	rs40125	rs37353	rs27184	rs35310	rs149108
rs2059175	rs35274	rs187645	rs1445951	rs26689	rs153980
rs2059174	rs244577	rs1809012	rs1947090	rs27187	rs153961
rs1118965	rs35267	rs187644	rs26708	rs1445948	rs1867725
rs154028	rs35266	rs153981	rs2112959	rs26687	rs153965
rs151802	rs39672	rs255652	rs1445953	rs166260	rs153966
rs244580	rs958851	rs255650	rs26709	rs149506	rs1988803
rs1457145	rs244576	rs255649	rs26710	rs27722	rs467300
rs244579	rs244575	rs2194210	rs28055	rs26695	rs1664886
rs255812	rs244573	rs255648	rs26711	rs27773	rs1867724
rs154029	rs35258	rs255647	rs27723	rs1471429	rs1445947
rs185333	rs35259	rs154221	rs27185	rs1471430	rs42470
rs35289	rs40121	rs256752	rs27695	rs26705	rs1423308
rs35288	rs35261	rs256120	rs1445954	rs28054	rs27174
rs35287	rs35264	rs255635	rs27549	rs26703	rs168834
rs35286	rs40122	rs185325	rs455969	rs27898	rs27727
rs35285	rs35265	rs26686	rs26712	rs722010	rs27172
rs35284	rs35255	rs1031197	rs1867711	rs27957	rs676449
rs35283	rs721826	rs1031198	rs1867712	rs26702	rs27186
rs35282	rs244570	rs27183	rs26713	rs27548	rs2112957
rs35281	rs27171	rs28044	rs26714	rs26701	rs1023814
rs35280	rs1824159	rs27182	rs27547	rs27188	rs27175
rs35279	rs27170	rs545611	rs26715	rs27189	rs1445950
rs35278	rs27169	rs649476	rs27949	rs149084	rs2021384
rs40126	rs27168	rs1664896	rs26700	rs153968	rs736736
rs35277	rs2013979	rs149106	rs1306348	rs464787	rs745813
rs35276	rs889231	rs1374028	rs35309	rs153978	rs889229

rs1077978	rs1353749	rs2055295
rs2081106	rs1391651	rs1391648
rs1559252	rs1391650	rs2055298
rs2054443	rs1391649	rs1472456
rs922437	rs1391652	rs1553114
rs922436	rs950446	rs1542842
rs922435	rs950447	rs1498611
rs922434	rs1498599	rs1532520
rs716908	rs1498601	
rs1971940	rs1498609	
rs1559251	rs1498608	
rs1345791	rs1553113	
rs1345792	rs1353748	
rs1345793	rs1498606	
rs1105577	rs1353747	
rs1960	rs1006431	
rs1824788	rs1948651	
rs1862563	rs1498605	
rs1551939	rs1498604	
rs1038080	rs1498603	
rs997421	rs1995166	
rs1014317	rs1498602	
rs2059191	rs1077183	
rs1551938	rs1078368	
rs1186170	rs1874857	
rs986067	rs1874858	
rs954740	rs1909294	
rs1363882	rs1546221	

Table 10

New SNP's identified by deCODE

Position in patent	Variation	AA Change Exon	1268007	A/G
732790	G/T		1268187	C/T
735966	C/A		1268553	A/G
736226	A/G		1272669	G/A
736516	C/T		1272910	A/G
850001	G/A		1273023	G/A
852776	A/C		1273220	A/G
853079	G/T		1273240	A/G
853575	C/A		1273543	C/T
856468	A/G		1288439	G/A
860845	A/G		1289730	T/A
870924	A/G		1290176	G/A
1027267	T/C		1293745	T/C
1027643	T/G		1344605	A/G
1027757	T/C		1344864	G/A
1028146	T/A		1345135	C/G
1037657	A/C		1345286	A/G
1044016	G/A		1346112	C/T
1044045	C/T		1352976	A/T
1254737	T/C		1354291	T/C
1254849	T/C		1354377	C/T
1255763	G/T		1354554	C/A
1257206	A/G		1354675	T/C
1258161	T/C		1355114	T/C

1355693	A/G	1575634	A/T	
1357081	A/G	1580088	G/A	
1362985	T/G	1581078	G/A	
1363021	C/T	1582418	T/A	
1363827	C/T	1584580	A/C	
1363911	G/A	1585955	G/T	
1364061	C/T	1590608	T/C	
1364066	T/A	1590677	A/G	
1367904	A/G	1590673	G/T	
1368193	T/C	1590837	G/A	
1368217	G/C	1590936	C/A	
1373349	C/T	1591011	G/A	
1373384	A/G	1591047	C/T	
1373415	T/C	1591306	C/A	Pro->Thr D1
1373979	T/G	1591583	T/C	
1376149	G/A	1594788	C/A	
1384931	A/C	1594994	G/A	
1385093	A/T	1601831	C/T	
1385107	G/A	1636902	T/C	
1385445	T/C	1638550	A/C	Lys->Thr exon 4
1391418	G/C	1640663	T/C	
1409210	C/A	1641954	C/T	
1414804	C/T	1641960	C/T	
1428284	T/C	1653881	G/A	
1431800	A/T	1655748	G/A	
1449904	A/T			
1574301	C/G			
1574615	C/T			

While this invention has been particularly shown and described with reference to preferred embodiments thereof, it will be understood by those skilled in the art that various changes in form and details may be made therein without departing from the spirit and scope of the invention as defined by the appended claims.

CLAIMS

What is claimed is:

1.   An isolated nucleic acid molecule comprising a phosphodiesterase 4D gene, or a
        fragment or variant thereof.

5   2.   The isolated nucleic acid molecule of Claim 1, wherein the phosphodiesterase
            4D gene has the nucleotide sequence of SEQ ID NO:1 which may optionally
            comprise at least one polymorphism as shown in Table 9, 10 or combination
            thereof.

3.   A nucleic acid encoding a polypeptide having an amino acid sequence selected
10   from the group consisting of SEQ ID NOs: 2-10, 12 or 14.

4.   An isolated nucleic acid molecule comprising a nucleotide sequence selected
        from the group consisting of SEQ ID NO: 1 which may optionally comprise at
        least one polymorphism as shown in Table 9, 10 or combination thereof, and the
        complement thereof.

15   5.   An isolated nucleic acid molecule which hybridizes under high stringency
            conditions to a nucleotide sequence selected from the group consisting of SEQ
            ID NO: 1 which may optionally comprise at least one polymorphism as shown
            in Table 9, 10 or combination thereof, and the complement thereof.

6.   An isolated nucleic acid molecule which hybridizes under high stringency conditions to a nucleotide sequence encoding an amino acid sequence selected from the group consisting of: SEQ ID NOs: 2-10, 12 or 14.

5   7.   A method for assaying the presence of a first nucleic acid molecule in a sample, comprising contacting said sample with a second nucleic acid molecule comprising a nucleotide sequence selected from the group consisting of SEQ ID NO: 1 which may optionally comprise at least one polymorphism as shown in Table 9, 10 or combination thereof, and the complement thereof, under high stringency conditions.

10   8.   A vector comprising an isolated nucleic acid molecule selected from the group consisting of: SEQ ID NO: 1, the complement of SEQ ID NO: 1 SEQ ID NOs: 2-10, 12 or 14, operatively linked to a regulatory sequence; wherein the nucleic acid molecule may optionally comprise at least one polymorphism as shown in Table 9, 10 or combination thereof.

15   9.   A recombinant host cell comprising the vector of Claim 8.

10.   A method for producing a polypeptide encoded by an isolated nucleic acid molecule, comprising culturing the recombinant host cell of Claim 9 under conditions suitable for expression of said nucleic acid molecule.

20   11.   An isolated polypeptide encoded by a phosphodiesterase 4D gene, or a fragment or variant of said polypeptide.

12.   The isolated polypeptide of Claim 11, wherein the phosphodiesterase 4D gene has the sequence of SEQ ID NO: 1 which may optionally comprise at least one

polymorphism as shown in Table 9, 10 or combination thereof, or the
complement thereof.

13.   The isolated polypeptide of Claim 11, wherein the polypeptide has an amino
       acid sequence selected from the group consisting of SEQ ID NOs: 2-10, 12 or
   5   14.

14.   An isolated polypeptide comprising an amino acid sequence which is greater
       than about 90 percent identical to an amino acid sequence selected from the
       group consisting of SEQ ID NOs: 2-10, 12 or 14.

15.   A fusion protein comprising an isolated polypeptide of Claim 11.

   10 16.   An antibody, or an antigen-binding fragment thereof, which selectively binds to
       a polypeptide of Claim 11.

17.   An antibody, or an antigen-binding fragment thereof, which selectively binds to
       an amino acid sequence selected from the group consisting of SEQ ID NOs: 2-
       10, 12 or 14, or to a fragment or variant of said amino acid sequence.

   15 18.   A method for assaying the presence of a polypeptide encoded by an isolated
       nucleic acid molecule according to Claim 1 in a sample, comprising contacting
       said sample with an antibody which specifically binds to the encoded
       polypeptide.

19.   A method of diagnosing a susceptibility to stroke in an individual, comprising
   20   detecting a polymorphism in phosphodiesterase 4D gene, wherein the presence
       of the polymorphism in the gene is indicative of a susceptibility to stroke.

20.   A method of diagnosing a susceptibility to stroke, comprising detecting an
       alteration in the expression or composition of a polypeptide encoded by
       phosphodiesterase 4D gene in a test sample, in comparison with the expression
       or composition of a polypeptide encoded by phosphodiesterase 4D gene in a

5   control sample, wherein the presence of an alteration in expression or
       composition of the polypeptide in the test sample is indicative of a susceptibility
       to stroke.

21.   The method of Claim 20, wherein the alteration in the expression or
       composition of a polypeptide encoded by phosphodiesterase 4D gene comprises

10   expression of a splicing variant polypeptide in a test sample that differs from a
       splicing variant polypeptide expressed in a control sample.

22.   A method of identifying an agent which alters activity of a polypeptide of Claim
       11, comprising:

       a)   contacting the polypeptide or a derivative or fragment

15   thereof, with an agent to be tested;

       b)   assessing the level of activity of the polypeptide or derivative or
                fragment thereof; and

       c)   comparing the level of activity with a level of activity of the
                polypeptide or active derivative or fragment thereof in the absence

20   of the agent,
       wherein if the level of activity of the polypeptide or derivative or fragment
       thereof in the presence of the agent differs, by an amount that is statistically
       significant, from the level in the absence of the agent, then the agent is an agent
       that alters activity of the polypeptide.

23.   An agent which alters activity of a polypeptide encoded by phosphodiesterase 4D gene, identifiable according to the method of Claim 22.

24.   An agent which alters activity of a polypeptide encoded by phosphodiesterase 4D gene, wherein the agent is selected from the group consisting of: a
5   phosphodiesterase 4D gene receptor; a phosphodiesterase 4D gene binding agent; a peptidomimetic; a fusion protein; a prodrug; an antibody; and a ribozyme.

25.   A method of altering activity of a polypeptide encoded by phosphodiesterase 4D gene, comprising contacting the polypeptide with an agent of Claim 24.

10   26.   A method of identifying an agent which alters interaction of the polypeptide of Claim 11 with a phosphodiesterase 4D gene binding agent, comprising:
        a)   contacting the polypeptide or a derivative or fragment thereof, the binding agent and with an agent to be tested;
        b)   assessing the interaction of the polypeptide or derivative or
15   fragment thereof with the binding agent; and
        c)   comparing the level of interaction with a level of interaction of the polypeptide or derivative or fragment thereof with the binding agent in the absence of the agent,
        wherein if the level of interaction of the polypeptide or derivative or fragment
20   thereof in the presence of the agent differs, by an amount that is statistically significant, from the level of interaction in the absence of the agent, then the agent is an agent that alters interaction of the polypeptide with the binding agent.

27.   An agent which alters interaction of a phosphodiesterase 4D gene polypeptide
25   with a phosphodiesterase 4D gene binding agent, identifiable according to the method of Claim 26.

28.   An agent which alters interaction of a phosphodiesterase 4D gene polypeptide
        with a first phosphodiesterase 4D gene binding agent, selected from the group
        consisting of: a phosphodiesterase 4D gene receptor; a second
        phosphodiesterase 4D gene binding agent; a peptidomimetic; a fusion protein; a

5   prodrug; an antibody; and a ribozyme.

29.   A method of altering interaction of a phosphodiesterase 4D gene polypeptide
        with a phosphodiesterase 4D gene binding agent, comprising contacting the
        phosphodiesterase 4D gene polypeptide and/or the phosphodiesterase 4D gene
        binding agent with an agent of Claim 28.

10   30.   A method of identifying an agent which alters expression of phosphodiesterase
        4D gene, comprising the steps of:
        a)   contacting a solution containing a nucleic acid of Claim 1
                or a derivative or fragment thereof with an agent to be
                tested;

15   b)   assessing the level of expression of the nucleic acid, derivative or
                fragment; and

        c)   comparing the level of expression with a level of expression of the
                nucleic acid, derivative or fragment in the absence of the agent,
        wherein if the level of expression of the nucleotide, derivative or fragment in the

20   presence of the agent differs, by an amount that is statistically significant, from
        the expression in the absence of the agent, then the agent is an agent that alters
        expression of phosphodiesterase 4D gene.

31.   An agent which alters expression of phosphodiesterase 4D gene, identifiable
        according to the method of Claim 30.

25   32.   A method of identifying an agent which alters expression of phosphodiesterase
        4D gene, comprising the steps of:

a)   contacting a solution containing a nucleic acid comprising

the promoter region of phosphodiesterase 4D gene operably

linked to a reporter gene, with an agent to be tested;

b)   assessing the level of expression of the reporter gene; and

5   c)   comparing the level of expression with a level of expression of the

reporter gene in the absence of the agent,

wherein if the level of expression of the reporter gene in the presence of the

agent differs, by an amount that is statistically significant, from the level of

expression in the absence of the agent, then the agent is an agent that alters

10   expression of phosphodiesterase 4D gene.

33.   An agent which alters expression of phosphodiesterase 4D gene, identifiable

according to the method of Claim 32.

34.   A method of identifying an agent which alters expression of phosphodiesterase

4D gene, comprising the steps of:

15   a)   contacting a solution containing a nucleic acid of Claim 1

or a derivative or fragment thereof with an agent to be

tested;

b)   assessing expression of the nucleic acid, derivative or fragment;

and

20   c)   comparing expression with expression of the nucleic acid,

derivative or fragment in the absence of the agent,

wherein if expression of the nucleotide, derivative or fragment in the presence

of the agent differs, by an amount that is statistically significant, from the

expression in the absence of the agent, then the agent is an agent that alters

25   expression of phosphodiesterase 4D gene.

35.   The method of Claim 34, wherein the expression of the nucleotide, derivative or

fragment in the presence of the agent comprises expression of one or more

splicing variant(s) that differ in kind or in quantity from the expression of one or more splicing variant(s) the absence of the agent.

36.   An agent which alters expression of phosphodiesterase 4D gene, identifiable according to the method of Claim 34.

5   37.   An agent which alters expression of phosphodiesterase 4D gene, selected from the group consisting of: antisense nucleic acid to phosphodiesterase 4D gene; a phosphodiesterase 4D gene polypeptide; a phosphodiesterase 4D gene receptor; a phosphodiesterase 4D gene binding agent; a peptidomimetic; a fusion protein; a prodrug thereof; an antibody; and a ribozyme.

10   38.   A method of altering expression of phosphodiesterase 4D gene, comprising contacting a cell containing phosphodiesterase 4D gene with an agent of Claim 37.

39.   A method of identifying a polypeptide which interacts with a phosphodiesterase 4D gene polypeptide, comprising employing a two yeast hybrid system using a

15   first vector which comprises a nucleic acid encoding a DNA binding domain and a phosphodiesterase 4D gene polypeptide, splicing variant, or fragment or derivative thereof, and a second vector which comprises a nucleic acid encoding a transcription activation domain and a nucleic acid encoding a test polypeptide, wherein if transcriptional activation occurs in the two yeast hybrid system, the

20   test polypeptide is a polypeptide which interacts with a phosphodiesterase 4D polypeptide.

40.   A phosphodiesterase 4D gene therapeutic agent selected from the group consisting of: a phosphodiesterase 4D gene or fragment or derivative thereof; a polypeptide encoded by phosphodiesterase 4D gene; a phosphodiesterase 4D

25   gene receptor; a phosphodiesterase 4D gene binding agent; a peptidomimetic; a

fusion protein; a prodrug; an antibody; an agent that alters phosphodiesterase 4D gene expression; an agent that alters activity of a polypeptide encoded by phosphodiesterase 4D gene; an agent that alters posttranscriptional processing of a polypeptide encoded by phosphodiesterase 4D gene; an agent that alters

5   interaction of a phosphodiesterase 4D gene with a phosphodiesterase 4D gene binding agent; an agent that alters transcription of splicing variants encoded by phosphodiesterase 4D gene; and a ribozyme.

41.   A pharmaceutical composition comprising a phosphodiesterase 4D gene therapeutic agent of Claim 40.

10   42.   The pharmaceutical composition of Claim 41, wherein the phosphodiesterase 4D gene therapeutic agent is an isolated nucleic acid molecule comprising a phosphodiesterase 4D gene or fragment or derivative thereof.

43.   The pharmaceutical composition of Claim 41, wherein the phosphodiesterase 4D gene therapeutic agent is a polypeptide encoded by the phosphodiesterase

15   4D gene.

44.   A method of treating stroke in an individual, comprising administering a phosphodiesterase 4D gene therapeutic agent to the individual, in a therapeutically effective amount.

45.   The method of Claim 44, wherein the phosphodiesterase 4D gene therapeutic

20   agent is a phosphodiesterase 4D gene agonist.

46.   The method of Claim 45 wherein the phosphodiesterase 4D gene therapeutic agent is a phosphodiesterase 4D gene antagonist.

47.   A transgenic animal comprising a nucleic acid selected from the group
        consisting of: an exogenous phosphodiesterase 4D gene and a nucleic acid
        encoding a phosphodiesterase 4D gene polypeptide.

48.   A method for assaying a sample for the presence of a phosphodiesterase 4D
5   gene nucleic acid, comprising:

        a)   contacting said sample with a nucleic acid comprising a contiguous
                  nucleotide sequence which is at least partially complementary to a
                  part of the sequence of said phosphodiesterase 4D gene nucleic
                  acid under conditions appropriate for hybridization, and
10   b)   assessing whether hybridization has occurred between a
                  phosphodiesterase 4D gene nucleic acid and said nucleic acid
                  comprising a contiguous nucleotide sequence which is at least
                  partially complementary to a part of the sequence of said
                  phosphodiesterase 4D gene nucleic acid.

15   49.   The method of Claim 48, wherein said nucleic acid comprising a contiguous
        nucleotide sequence is completely complementary to a part of the sequence of
        said phosphodiesterase 4D gene nucleic acid.

50.   The method of Claim 48, comprising amplification of at least part of said
        phosphodiesterase 4D gene nucleic acid.

20   51.   The method of Claim 48, wherein said contiguous nucleotide sequence is 100 or
        fewer nucleotides in length and is either: a) at least 80% identical to a
        contiguous sequence of nucleotides in SEQ ID NO: 1 which may optionally
        comprise at least one polymorphism as shown in Table 9, 10 or combination
        thereof; b) at least 80% identical to the complement of a contiguous sequence of
25   nucleotides in SEQ ID NO: 1 which may optionally comprise at least one

polymorphism as shown in Table 9, 10 or combination thereof; or c) capable of selectively hybridizing to said phosphodiesterase 4D gene nucleic acid.

52.　A reagent for assaying a sample for the presence of a phosphodiesterase 4D gene nucleic acid, said reagent comprising a nucleic acid comprising a contiguous nucleotide sequence which is at least partially complementary to a part of the nucleotide sequence of said phosphodiesterase 4D gene nucleic acid.

53.　The reagent of Claim 52, wherein the nucleic acid comprises a contiguous nucleotide sequence which is completely complementary to a part of the nucleotide sequence of said phosphodiesterase 4D gene nucleic acid.

54.　A reagent kit for assaying a sample for the presence of a phosphodiesterase 4D gene nucleic acid, comprising in separate containers:

a)　one or more labeled nucleic acids comprising a contiguous nucleotide sequence which is at least partially complementary to a part of the nucleotide sequence of said phosphodiesterase 4D gene nucleic acid, and

b)　reagents for detection of said label.

55.　The reagent kit of Claim 54, wherein the labeled nucleic acid comprises a contiguous nucleotide sequences which is completely complementary to a part of the nucleotide sequence of said phosphodiesterase 4D gene nucleic acid.

56.　A reagent kit for assaying a sample for the presence of a phosphodiesterase 4D gene nucleic acid, comprising one or more nucleic acids comprising a contiguous nucleotide sequence which is at least partially complementary to a part of the nucleotide sequence of said phosphodiesterase 4D gene nucleic acid, and which is capable of acting as a primer for said phosphodiesterase 4D gene nucleic acid when maintained under conditions for primer extension.

57. The use of a nucleic acid which is 100 or fewer nucleotides in length and which is either: a) at least 80% identical to a contiguous sequence of nucleotides in SEQ ID NO: 1 which may optionally comprise at least one polymorphism as shown in Table 9, 10 or combination thereof; b) at least 80% identical to the

5 complement of a contiguous sequence of nucleotides in SEQ ID NO: 1 which may optionally comprise at least one polymorphism as shown in Table 9, 10 or combination thereof; or c) capable of selectively hybridizing to said phosphodiesterase 4D gene nucleic acid, for assaying a sample for the presence of a phosphodiesterase 4D gene nucleic acid.

10 58. The use of a nucleic acid which is 100 or fewer nucleotides in length and which is either: a) at least 80% identical to a contiguous sequence of nucleotides in SEQ ID NO: 1; b) at least 80% identical to the complement of a contiguous sequence of nucleotides in SEQ ID NO: 1 which may optionally comprise at least one polymorphism as shown in Table 9, 10 or combination thereof; or c)

15 capable of selectively hybridizing to said phosphodiesterase 4D gene nucleic acid, for assaying a sample for the presence of a phosphodiesterase 4D gene nucleic acid that has at least one nucleotide difference from SEQ ID NO: 1 which may optionally comprise at least one polymorphism as shown in Table 9, 10 or combination thereof.

20 59. The use of a nucleic acid which is 100 or fewer nucleotides in length and which is either: a) at least 80% identical to a contiguous sequence of nucleotides in SEQ ID NO: 1 which may optionally comprise at least one polymorphism as shown in Table 9, 10 or combination thereof; b) at least 80% identical to the complement of a contiguous sequence of nucleotides in SEQ ID NO: 1 which

25 may optionally comprise at least one polymorphism as shown in Table 9, 10 or combination thereof; or c) capable of selectively hybridizing to said phosphodiesterase 4D gene nucleic acid, for diagnosing a susceptibility to stroke.

Fig. 1A

Fig. 1B

Genetic map   Combined map - cM   Physical map - Mb

Fig. 2A

D5S1968

D5S474

D5S491
D5S2102

D5S1715

D5S2028

D5S2107

D5S2091

D5S2080

D5S624

D5S1990

D5S1474

D5S427

D5S668
D5S2858

D5S2072
D5S2089

D5S647

D5S2048

D5S2046

Fig. 2B

**D5S1968

2.26

D5S474
*D5S664
*D5S633   1.97
*D5S461
*D5S407
D5S491
D5S2102
1.47

D5S1715
*D5S398   0.93
D5S2028
0.576
D5S2107
*D5S468   0.85
D5S2091, *D5S2000
0.77
D5S2080
*D5S2018   0.59
*D5S2071
D5S624
1.15
D5S1990
D5S1474
D5S427   0.4
*D5S1956   0.2
D5S668
D5S2858   1.49
D5S2072, D5S2089
0.58
D5S647
0.42
D5S2048, *D5S2121
0.49
D5S2046

*Markers only assigned in physical map
**Marker in blue - only assigned in genetic map

Fig. 2C

D5S474
D5S664

2.6 Mb   D5S633   0.46
D5S461   0.1
D5S407   0.8
D5S491
D5S2102   0.1
0.1
0.4
0.4
0.2

D5S1715   0.6
D5S398   0.1
D5S2028   0.5
4.3 Mb   D5S2107   0.2
0.6
D5S468
D5S2091, D5S2000   0.1
0.4
D5S2080   0.07
D5S2018   0.08
D5S2071   1.5
D5S624   0.2

2Mb   D5S1990   0.4
D5S1474   0.3

D5S427   0.5
D5S1956   0.1
?
D5S668
6.7 Mb   D5S2858   0.2

appr.
1.6

D5S2072, D5S2089
D5S647   0.2
D5S2048, D5S2121   0.5
D5S2046   0.5

1.2

Fig. 3

Fig. 4

5/.375

Fig. 5

>Contig_2 (1,1691140)
CATTTTTTGAAAGAACATCTGAAAGACAAAATGGGGAATGGCGAGTCTGTCTAATAAACCATTTTGAGAAAACTGGATA
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CTTAATATCAGAAATGATGAAACTTTTAAAAGAAAACATAGGGTGAATGTATGTTTAGGGAAACTCAAATTTATGTTCA
CAGGTTGCAAAAAGAAAAATATGAGAAAAACATAGAGGAAAATGATTCTGCCAATAAAGTGAGTTGGAAATAATTTTTC
TGTTTTCACAAAAATCATTGCTAACAAAAGCAAAAACAAGTGTGGGACTATAGAAACTGATGAGCTTCTGCATAGGAA
AGAAAAGAATGAACCAATACAGAAGGCATCCAGTAGATTGGGACAAAATTATGGGGGATTATATATCTGAAAGGGTTGT
TATCTAACATGTATAAGAAATTACCACTACTAAGTAGGAAAAACAACAACACAAAACAACAAATAACCAGATGGAAATT
GGGCAAAGAACCTGAATAGATGTTTCTGAATAGAAGACATGAATTTGACTACCAGGTAAAAGAAAAGGTTCTCAACATA
CCTAATCATCAAGAAAATGTACATTAAAACTCACTGAGATATCTTCTCCACTCTAATTGGAATTAATGTTACAAAAAAG
AAACAAATTTTTTACAATGAAATGATCAGTGTGGAGTTGGATGAAGGGGTACTATTACACTACTACACAGTGTAGGGTG
GAATTTAAGTCAGTATACACACTATGAAAAATAGTTGGAGTTTGCTCAAAAAAATAAAATACAACTATCATTTGCTGTAG
TAATCCCACCACTGAATATACATTTTAAGAAAATGAAATCAGTATATTGAAGAGATACGTGAAATCCTACATTTCTTGA
CACATTATTCACAGCACTCAAGATGTGGAATCAACCTACCTGTCCAGGAACAGATGAAGAGATAAAGAAAATGCAGTGT
GTATACACAGTGGAATGCTCTTCACCATAAAAAATTCACGGAATCATGTCATTGCAGCAACATGGTGGACAATGTAAGA
AAAGCTCCCCGGAGAAGCTGTACAGAAGCTGCCTCCTCAGCAGTCAGGGCCAGGGACCGGAGCTGTTTTTACCCCAGGA
CAGGGCCGGCCCCAAGTCATCCCAGAGCTGCCATGGCACCCCCTCAGTCGGGTCCTGAGGAATCCTACACAAGCTACTT
ATATCAGTGATCACTAGGATAATCCATAGAACTTTTGGGAAAGAAGTTTAAGACCTTTCTCCCACCATTTCAGCAGGAT
AAATTCCAACTGGATTAGAAAATGAAATGTTAATAATGCAAATAAGTACATATTTATATCTGTATATAAAATACAGTTG
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ATTCACTTATTCCACTTCAAGCCAGCTAAATGATTGTTTCCCTGATGGCAAAAGTCTCAGATTGATTGCACAGTTTATT
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CAAGAAACAAGTTACCTTTGGGGCAACAGAGGCCCTTGTCATTTTCTCAAAAGAAGGAAGCATCAGCATTTTGATGATG
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GAACCTTAAAAAAAAAGAGAAGCTTTTAAAAAAATCATAATAGTTTATGATCTTGAAGGGTTTAAAAGTATTTGATGAAGA
TGTCTTTTGAATTTATTTGTAGGTCTTCTTGTGTATTTAAAAGCTAAGTTATCTTGTAATCATTTTTTTCTATACCTTT
GTCAGTAACCTCTTAGTGATGAAATAAAAAAGATTAGGTAATCATCCAGCAATGGGGAAGAAGTTAAGGAACAAAGAGC
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CCTCATAGATCACCCATGTGCTGAAGACTTTCAGTTCTGTATCTTCATTCTAGATCTCCTGAACTCAAGATCAGAATAT
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AGTGCATTGTATGTAGCAAGAGTAGGTATTTTCGTAAACTTTTTGTTTTAATTAAACACATATATATTTATTATTGCAG
TGTAAAATGTATTTCTTAATTTGGATAAGTGTTAGTGAGGATGTGGATAAATTGGAACTCTTGTACATTACTGGTGGGA
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ACTCCAGAAGTTAAAGATCAGTGAGCTTGAATACACACAATAGAAGCTAGTCTAAACAAAGCACAGAGAGAAAAAAGAA
CAAAACAAACCTCCCACAACAAAACAAAACAAAAGCCAACCAAAAATACAGCCTCAATGACCTATGGGAAATATTTAGC
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TTAAGATTTTATTATTGATTTTTAGGAATTGCATTATATCTTGGTGTGGTTGTTTAAACAGAGGTATAGCTTATCAACC

Fig. 6.1

AATGGTGGAGCTAAAATAGAATACTTGAAAGTACT.ATGGATGCACAGAATCTAAGATGGCCCCCAATTTTCCTGCTAC
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CAAGGGTGACATGTAATTAAGCAAAGCTCAGTAAATTTAAAATGATTGAAATTGTACTAAGTTTTCTGACCACGCTAGA
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GATTGCAGAAAATGTATGTGACAATATTCAACACCCATTAATGATAAAAATGTTAAATACATTACAATAGAAGAAAACT
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CTACAAAGGACATGAACTCATCCTTTTTATGGCTAGCTACTAGTATTCCATGGTGTATGTGTGCCACATTTTATTTTTATT
TTTATTATTTTTATTTTTAAAATTTTATTATTATACTTTAAGTTAGTGTACATGTGCACAACATGCAGGTTTGTTACAT
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CCCACCTATGAGTGGCAACATGTGGTGTTTGGTTTTTTGTCCTTGAGATAGTTTGCTGAGAATGATGGTTTCCAGTTTC
ATCCATGTCCCTACAAAGCACATGAACTCATTATTTTTCATGGCTGCATAGTATTCCGTGGTGTATAGTGCCACATTTT
CTTAATCCAGTCTATCACTGATGGACATTTGGGTTGGTTCCAAGTCTTTGCTATTGTGAATAGTGCCTCAATAAACATA

Fig. 6.2

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CGTGTGCATGTGTCTTTATAGCAGCATGATTTATAATCCTTTGTGTATATACCCAGTAATGGGATGGCTGGGTCAAATG
GTATTTCTAGTTCTAGATCCTTGAGGAATCGCCACACTGTCTTCCACAATGGTTGAACCAGTTTACAGTCCCACCAACA
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TTGTATAAGGTGTAAGGAAGGGATCTAGTTTCAGCTTTCTACATATGGCTAGCCAGTTTTCCCAGCACCATTTATTAAA
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GATTATGGTGGATCAGCTTTTTGATGTGCTGCTGGATTCGGTTTGCCAGTACTTTATTGAGGATTTGTTCATTGATGTA
CATCAGGGATATTAGTGTAAAATTCTCTTTTTTTGTTGTGTCTCTGCCAGGCTTTGGTATCAGGATGATGCTGGCCTCA
TCAAATGAGTTAGGGAGGATTCCCTCTTTTTCTATTGATTGGAATAGTTTCAGAAGGAATGGTACCAGCTCCTCCTTGT
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CCTCGATGGTCTTTACAATTTGGCATGATTTTGCAGTGGCTGGTACCGTATGTTCCTTTTCCATGTTTAGTGCTTCCTTC
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CACTTATGAAGCTTAGTTTGGCTGGATATGAAATTCTGGGTTGAAAATTCTTTCCTTCAGGAATGTTGAATATTGGTCC
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AGCTGCGTTCCTTTGGAGGAGGAGAGGTGCTCTGATTTTTAGAGTTTCCAGTTTTTCTGCTCTGTTTTTCCCCCATCTT
TGTGGTTTTATTTACCTTTGGTCTTTGATGATGGTGACGTACAGATGGGGTTTTGGTGTGGATGTCCTTTCTGTTTGTT
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Fig. 6.3

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AGTTTTCCTTCTAACAGTTAGGACCCTCAGCTGCAGGTCTGTTGGTGTTTGCTGGAGGTCCACTCCAGACCCTGTTTGC
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GGGAGTCAGGGACCCACTTGAGGAGGCAGTCTGTCCATTCTCAGATCTCAAGCTGCATGCTGGGAGAACCACTACTCTC
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Fig. 6.4
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Fig. 6.5

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Fig. 6.6

ATAATAGTAGTTAATATTTACCTCATAGAGTTTTTCTGGGGAATTAAAACATGTGATTAAAATACATGTGGGCTTAATA
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CACTATCAAATTAGTTTCTTGCCTCAGGTATCAGTACAAGTTTTTAGTTCCTCTTATGTTAGGGGAACACAGCTTAGGG
TTCCAGCGAATCGTTTTTAACTCACAATATGTTACTTTTGACTATTCATAAAGGTTGATCCATGATCAGGTGAGTGGTT
TTTGTTTTTAAAGACTGAACTCTACATAGTTGGATTCAGGGAACACCGTACATGTGGGATTACACACAGAGCACCAGAG
TTCTGGGCTTCTGATCATATTATTCTCAGAGTAATTTTGGAGGAGAGTTTAGTGGTAAAGATCAACAGAAAGTATACTG
CCTCTCTTTTAAGCTTTTCACAAAGTAATATACACCAGAATATTCCCTCTTCCATTTGAGCAGAAGCTGGTGGGGCTGA
GTTGGAGAAACAGAGAAATGATGAAATGTAATTGGGCCAAAATTTTATTTAACTGTGTAGAACAGAAGTTATTTTGAAG
TCAGAGTTACTTGTGAATATATAAAAAGTGAGGTCTATTTATAGTTGGCAGTGGGGGGAATTTTCAGAAACATGATTAA
AAGGGGAACTTATTTGAAGCTTAGATCAAAGGAAAGATGGATTAGCTCTATAGAAATTGGAATGAAATGTTTTAACAAA
AATAAACAAAAACAGGAAAATACAGGCTCAGAAACTCTTAAAAATTTGGCTGCCAAATTTTATTGGCTTTTATGTGTCTT
TTTTTACTCTTCTTGGACCATTCATTTTTTTAGTTTCAGAAATCCTATCCTGAAGTCTATCAAGAACGATTTCCTACAC
CAGAAAGTGCAGCACTTCTGTTTCCTGAAAAACCCAAACCGAAACCACAGCTGCTAATGTGGGCACTAAAGAAGCCTTT
CCAACCATTTCAAAGAACTAGAAGTTTTCGAATGTAATAATACTTCCACAGCAACAGGTGCTAGAGACCACTGTTGTTG
TTTTGAGTGAATGGTGGTTAGGAGAAAGACTTTGGTGGTGGAAGAAAGAAAAGCATAAAACAAAGACTACTGAAATATA
GATAAAGATTGCCTTAGTTTTTAAAAATGTTTGGCCATTAGTATTTTTATAAAACTCAATGCTAGTTTTAAAGTGTATA
AATTGGTTAAAAATTTATGAGTCAAATATATAGTGATAATGTTAACATGTTTGTAATTGCTACAGAATTTAAGGGTATTT
TTATCTCTGTGCTTTCTTTTTCATGGTGTTTATTAAATAATTGTGTATATACATCCTAGCTACTGATATCTTTATTATA
GCCTTAAGACTTAATTTTAAGTCTTAAAAATAGCGTGTATACTTGAATAAGAAAGCACTGGGTACTGTTACTGTGATG
CTATTGACTTAGTAGCCAATTATCATTTCTCCTGTATAAATTCCAGTTTTTATTGCTGCACATAAATTTTTTAATGTCT
TATATTGTGATAGCTATGTCTTTTATTGCAGATTTATTGGATGTTATGACAGATTTTACTAAAGCTAGTGTTTTTATAA

Fig. 6.7

CATATATATTAGTTGATGTTTACCTATAAGTGGAGTAGATTTTCATCTGCCTGCAATGGTATAATTTCAGTCTTAGCTA
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AAAATGAGACTACTTCCTTTATAAATATATTTTCATCCTTTTGAAAGTAAGTGAAATGTAAATAAACTTATTTTTTTTA
AAAATGCCATACCTCGTGTTTCTTTTTTGGCATTTATTTAGATCAGTTAGAGAGCATCCTCAAATGCCTTTGCCAATGG
TCTTCATCAGTCATTTATTTCTATGAGAAAATGAAAAGTAAGATTTGGGCTTTTGTACTGACTGGTCTTTAGGTGAGCA
CTTTGCCAATTTTTTAGCATTGCTTTGTCCTTTTGTAAAATTGTGGTGCACTTGGATTATGGAGCATCTGAAAGTCTTC
CTCTGAGCGCAATTACCCAGAAAGCTGAGAGCAACACCCACTAATTGGCTTACAGTACTCCTTGGACTGATTTTTCCTT
GCACTTAAGATTTGCTCCTGTTCTCTTGTAGCAAAATACCTCCTACATCTGATGCAGATTTTGCTTTTAAAAATGGACC
AAAGTATTCCTATTGGTTTGGGTACCACCTTACATTCCAAATACATAATGTAATGGGAGATTTTTAGTGTTTCAGGATT
CATTCTCAGACTTTGGCATTCTGTTTAGAGCCAAGAATAATTCTTCTTTCTGCTCTATTTTCACTCCGAGGTAGAGTTT
ATTTTTCAGAATCTCTTTTCCTGTTGCTTGTCTTTGGAAGTATATGTGGTTGTTTCTTTCCTTTTTTTTTAAGATTA
CTTTTTTAAAAAAATATAACATATTTATGGAGAGATCATAAAACATAAGCACCAATTAAGTGTGAAATTGATGAATAGA
TAATAGCAGTTATCCAAAGTGGTTGTCCCAATTAACGCTCCCATTAACAGTGTATGAGAACTCCTTTTGTTTAATATCC
TCACCGAAACTTGCTATCGTCAGAGTTTGAAATGTTGTCAATGTGGTGGTTATTAATAGTGTCTGATTATGGTTTTATT
TTGCATTTCCATGATTATTAATATGGCTGAACATTTTTTTCCATTTTTTTGACTTTTTCGTCTTTGTACAAATTTTGGT
ATCAGATTGTCAAGCCTNCATACACACACACACACACATGCAGAGACACACACTCACCCTGTTGAGTTTTTTTTTTTAATT
TTGATTGTATTATATGTATAAATCAATTTGCAGAGAATTGCTGTCTACAATATTAGGTCTTCTAGTCCATGACCATGGT
TTACCTTTCCATTTTATTTGAAATAAATGAGAAAAAAAGTTGCTTTTTTCTGTGAAGATTTACAAATCTTTNGTTAGAT
TTGTTCCCAAGTATTTGATACTTTAAAAAAAGTATCAAAATTTTCAAAGACTGTAGTACCCATTTCCTGTTTCAACAGT
TACTCAGCTGATCAATGTTGACGTTTAATTCTTTTGTAAATGCTGGTGTTGTAAATTTCCTTATAGTGTTTGTTGCTTC
TGTATGGTATAAAACTGATTTTTCCTGAGTTTTGTTTACAGTGAACCTTGCTAGACTTGTTAATTCTAAACATTTAGTA
TGATTCTTTGGCATTGTCTACNTATTAGGAAAAAAACAGCTTTTGCTTTTCCGTTCCATGGTAGACAGCATCTATATTG
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ACATTTCTAAATAATAGAATATATCATCAGTGATGGGGTATCACTTCTCAGATTAGATTCCAAAACTCTGGCTTCCATC
TTGCTTGTCTTCCTTCTTATTCTCTCCCTGGCTTGCTCTGAGGAAAGCCAGCTGTCATGTTGTGAGCTGCCCTTTCAGA
GGCCCACCAGGCAAGGAACTGATGTCTCTGGCCAACAGCCAGCGTGGACATGAAGCCTGCCAACAGCTTTACGAGTGAT
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TCCCTTCCCTTCCGTCCCTTCCCTCCCTTCTCTCCCTTCCCTTCCCTTCCTTTCCTCCCTTCCCTCCCTTCCCTCCCTT
CACTTTCCTTCTTTTCTTTCCTTTCCTTCCTTTCCTTCCTTCCTTTCCCTGGCTCCCTCCATCCTTTCCTTCTTTTTTG
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CCTGGTAGACCAGGTACATTGTNAATGAGCAGTAATATTTTGAAAGGAATCTTTTTTTTTTTTTTTTTTTTTGAGCA
GCAGATCTCAATAGTGGGCTTAAAATATTGAGTAAACTATGCTGTAAACAGATGTACTGTTATCCAGGCTTTGCAGAGC
ACAGGCAGAGTAGATATAGTGTATTTTTTAAGGGCCCAAGGATTTTTTGAATGATAGATGAGCATTAGCTTCAGCTTCA
GGTCACCAGCTGCTAACAGGAGAGTCAGCTTGTCCTTTGAAGTTTTGAAGCCAGGCATTGGCTTCTTTATAGCTATGAA
AGTCCTAGATGACATCTTCTGCCAATAGAACGCTGTTTTATCTACAGTGAAAAGCTATTGTTAGTGCAACCACCTTCA
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CTCTCTCAGTCTTTATAGAATTGAGGAGTTAGGGCCTTGCTCTATAGTAGGCTTTGGCTTAAGGGAATATTGTCGCTGG
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CTGGAGTGGTGCTTTTAATTTCCTTCAAGAACTATTCCTTTCCACTCACAACTTGACTAACTGTTTGGCACAAGAGACC
TAAACTTTCCTTCTATCTTGGCTTTCAACATACCTTTCTCATTAAGCTCAATCATTTATAGCTTTCGATTTAAAGTGAG
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TGGGAATAGTGAAGCCCAAGGAGAGGGAGAGTGATGGGGGAATGGCTGGTGGGTAGAGCAGTCAGAACACACAACATTT
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GGAATAGAGTGGGGATCGCAGCAGCATCATCTTGCTCCTAATTTCAAAGAGGAGGTTTTTAGCATTAGAGTATTTGTAG
ATACCCTTTTATAATTTTAAAAGAAGTTATTTTCTATTTCTATTCTATTGTCAAAAAATTTTTATTGCTCGTTTTAACC
ATAAGTTGATGTTGAATCTTAATCAACTACCTTTCCTACATTTGAGGATTTTATAAGTTCTATCTTTTAGTCTATCATT
GGGGTTATATTACATTAATTGATTACTAATATTAAGCCACTTTGCATTCTAGGAGTGGCATAAATCTAATTATGATGTA

Fig. 6.8

TTATCATTTGAATATATATATCTAGATTCTGTTGCTAATATTTAGTTTATAGTTCTTGTATCTATGTTCATGAGTAAAA
TTGGCCTGAAAATTTCCTTCTCTTACTATCTTTGTTGACTTTCTGATCAAGGTTATAAATTAAGCTGAAAATTACTCTC
TTTTTCTATATATGGTTGGAATTATTTCTTTCTTGAATCTTGGGTGAAACTTAGTTAACTTTTAAAAAATAAATAATGT
AATTACACCGAAGTCAGAAAAATGGTTTGTATAATTTCAAGTCTTTGAAATTTGTTGAGACTTGTTGCATAGCCTAGTA
TATGGTCAATTTTTGTTAATATTCTGTACACATGAAAAGAATATGAATTTTGCATTGATTACAGTTTTTTTTCTCTCTCT
ATAAAATCTTGTTTAAATTTCCTACAATTTTACTGATTTATTTTGCCTGCATATATATGTGTGTGTGTGTGTGTGTGTG
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CTGGAGTGCAGTGGTGCAGTCATAGCTCACAGCAGCCTTGGCCTCCTGGGCTCAAGTGATGCTCCTGTCTCAGCCTCCC
AAGTAGCTAGGACTAATTATGCTCAGCTAATTTTATTCAAATTTTTTAGAGATTTTGTCTTGCTGTGCTTGTTACCCAG
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ATTTCTGATTAAATTCATTTTGAATTTTCATATTTATCTGGTGGACTGACCTTTTTATTTATTTATCATTATAAAATGTCATT
CTTTATCATTAGTAATGTGTTTGTCTGTAGTCTTTCTGTATTATATTTTAGGTGTATCTCTTATAAACAGCATGTAGTT
TATGTTTTAAGAAGCCAGCTTCACAGTCTTTATTTTTAATTGTAGCATTTAGACAGTTTATATTTAATACGATTACTCT
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CCTTTGTTTGGATTTTGGAAATAATTTCATGGTTTCTCCTGCTAATTTTGAAACTTATGCACTGTTTCCAGTCTTCTTG
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AAACTTCATAATATATTGTTGTTATTTTATATAGTCTTTGTTTACTAGACTTACCCATGCATTTTACCACTTTTTTTCT
TCTTAAAGTAAATTCTTTAGAATTTATCTGCAGTAAGGGATTGCTTTGTTTTTGTCTGAAAACTTTTTAAATTTTGCCC
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TTGTTTTATTTTTAGGTATGATTTTTATAGTATTTTTCATTTTTATGTATAATTGCTTTTATTTTTAGGTATGATAAGT
CAAAAATGTCATAGGGAGTATCATTGCCCTAAGAGACTCTGGTAAGAAATTTATCAAATGAATGTTCCATACAGAAAAT
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GCTGTGGTAAATCATTACTTTTACCCTCAAGAACAAAACCCTATATATGTCAAAGACCTAGGGAAAAGTAAGAGTTTTA
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TGGGACAACAGGGTGCTAACCCAATAAAAAAAGTCACTTTCAGTTGTTAGTGCATATTTATGTTGCAATGTAAGTTTC
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GTATCTAGAGCCTTTTCTAGGGAATATAACAAAAGCTTTTCTCCCTCTTCTAGCCCCCACAAATTTTACATCTTGCTGC
CAGCGTATCTTTCAAAATAATAACAAGGTAGTCTTCTTGAGTAGTTTATAATGGATTTCTTTCTATGTTTAATAAAGCA
AATGAAACTTAATTTTTTAAAAAAACTAGAGGACTAAAAATTATTTTCATGAGAACTAAGTAAAATATAATTGATATTTG
TATTAGTCCATTCTCATGGTACTATGAAGAAATACCTGAGACTGGGAGATTCATAAAGAGAAGAGGTTTAACTGACTCA

Fig. 6.9

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CAGTTCTGCATGGCTGGGGAGGCCTTAGGAAACTTACAATCATGGTGGAAGGTGAAGCAAACACATCCTTTTTCACAAG
GCGACAGGAGAAGAAGCGCCGAGCAAAGGGGGAANAGCCCCTTATATAACCATCAGATCTTGTGAGAACCCACTATCAT
GAGAACAGCATGGGGATAACTGCCCCCATGATTTGATTACCTCCCACTGGGCTCCTCTCACGACACGTGGGAATTATGG
GAACTACAACTCAAGATGAGATTTGGGTGGGGACACAGCCAACCACATCAATATTTAAAATTGAAGTACTTCAGTTTTG
ATTAGTGTAGACTAATACAACATGCGAATGTAGAGGCCTTTTGCTCACTCTTCCCACTAAGAACTCAAGACCTTTGAAG
GGTCCTGACAACAGACATTTCAGTGATAATTTACCAAGATAGCAGAACTGTTTGTTGCTAAAAATAGTTATGCAGGATA
AAATGCAATCTATAATTTTATGAGAACCTGTGATTCTAGTTTACCTCCAATTACTAATATCCATATCTATGTTTGTATA
TTTTTTAATCAGGTGTGTTTTGGACTCAGAAGTTATTAAAAAATTCAGTCAAATGATGGGAGTGATGTTTTGAAGATTAG
AATCATGTAGCTGATTCACATGCGTTATTAGACCATGATAAGAAGCACAGAGGAAATATAACAAATGAAATCAGCCTTC
TTGCCTACTGATTCTGCATAATTTTAAAAATTAAATTTGGGTTTTTGTCCATATGCAAATTTGTCCTAAAAATTTCCAT
CTTTTGAAAGCGTTTTGATACTTAGAAAAAAAAATTTAAGACTTTTCTAAATAGAATATACAAGATGGTACAATCTGAT
TGGATAATAATGCATCAGTGTTCTAAGTTAATAATTGTCTAAGGCAATAATTAACATTTAAAACATGCAGTGACATCAT
TTCTGAATTTTTTTCTACTTTCTACTTATCAAAATGCCAAATATAATTATACACAATTACTGCTTAAAGATGATGAAGG
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GTGTTGATAGTTATTGTATCATAAAGGAAAAACTAGGGATTTTATGTTTAAGTCCAGTATATTTGATATAGGTAGTAAG
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CCTTTATTACCATATGTTTTCCTCAAAACCCCAGAAAGCTTAACTGCTTGCTAAATACTCCCTAAAAGGAACTTACCAA
TGGTAGAAATGAACAATTATCTAAACGTAGCAACAGATAATTTGGGATAAATGTCAGATTATTGGATTTGAAACTGAAA
ATATTTCTTGCAAATTTCAGTTTGAGAAGAGAAACACAGAGGTAGGATAATTTCACGCATCAGCCCCTCACCTTAGCTG
TCTTCTCTGCAATATGGGGACAACTGTGCTTGTTCTCTTTCCCTCAGATAAATCTTGTTTTTATTCCTCAGACTAATGTT
TTGGTCTATAAACCGAGAATTTTATAATAAATGCGGAGATGGCAATCGGAGATAAAGCATATGTTTACAAGTAATGCCT
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TAAATGACTTTACTGCAGGTAGATGGTGTGATGGAAAGAGCAGTGTGTTAGCAGCAGTGGTTCTGCCACCAGATAGCTT
GAGGAGGCTTTAATTTCCCACACCTTAAATAAACATAATTAAGTAAACTGTGTTTGCTATGCAAATACAGTCTAAAAGC
AAGACTCCTAGTTCCTTCTTTCGAACTAAATTATTACCCAACGGTCCTTCCTATTAGAATTTGTAGCACCAATATTAGT
TAGAAATCAGAATATTTGGATTTTAACTTGTGGCACTGCCAGTTGGGCAAGTGACCTTGAGCAAGTCACAACCTCTCTG
AGATTTGTCTTTTTCTCAGTAAAGTGAAGGGATTGGATATGGAAGGAACAAACTCCAGTGTTTTCCAGGACAGGAAGAT
ATTAGAAGTTGCTGAGAATGGAGGGGTCAGTGGAGAAATGGAGTGTGCAGACCTTACAATAAGGTATCTGATAGAAACA
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GTTTCCTGAGCATGAGCTTTGTAGGCGACACATTTCAGGAGCTGAAACAGCCCAGACCCCACCTCTCTTTTCAGGCTCC
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GTTACTGAGGAGCTAATATTCAAGCTGAACCCTGAAGAATGTGTCCACTAGTGAGGAAGGCAGGGCAGAGGCATTTTCT
CAGAGGAAATAGCATGAAATAGGATCTGGTGACAACTAGAAATGACTTACCATTCCATGTGGTCCAGGTAATGCTAATA
CTTCAATATAAAGTCTGACATGGCTCTCTGTAAAGGACTGAGTAATAAGGGGATTTCATGAACTGTGGCCAGGCAGGCA
GCGGACTTATTTTTAAGTAAGGAGTTTTTTGGCACCTGTCAAATAACATCTCCGAGATGGCTTCCTTTACATATTCCAG
GTCAGTTGATCCAGATCTCAATGCTCTTTGCAATTTTTGGATTCTTAGTAGGTTGCTGCTACACCATGGTTTCTCTTTC
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GTTGCCTGCAGTGAGATGCTTTTAATATTTTCTCCACATGTGAACTCAGTGTTTTCCTTTAGGTTTTCTTTAATTTCCA
TGTGGTTAATGTCACTACTGCAGGGAGCCTGGATATTTATAGATTTTTCCCTTTCACTCTATCTTCAATAATTTGCTTT
TGCATTCAGTGAACACATCCATCACAATAAATGGTCCTTTTCTCATTTGTATTGACTTGCATTTATATAAACCAGACTT
CCTTTTTATAGTATTATCAATGTTCATGGCAATTTTCAGTAGCTTGGGGAGCTGGGTATGAGAGTTGGCACAGAGCTGC
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TTCCTTAATTGCCTCAATATAAAATGAGAGTATTAAATAAGCATCTCTAAATTATTCCCAATTTCATTAAAATGTTCTT
GATTTGATAGCCCTTTTGTGCACTGAAAGATGTAAAATATTCCAATTACATTGTTGACTATTCATAGTCAGTTCAATGT
TATAAACTTGTATAGCTCCTAGTAGAAATAAACAGGGAACTGAGTTATATACAGTTCATATTGATCATTCTTAGATGCT
ATTGCTTTCTGATAGCTCAATTTATTTTAACAGAGTCTACAATGATGTCATTTTAAAATTATTATTTTAATATTTCATT
TATATTCTTTTAATATAAGTTAATTTTTTTCCTATCACTGTGCATGTCATTGGTTTGCAGATTGTTGGGAAGAATATCAC
TGGCAGGACTTGTATTGCTGTTCTTTAGGCAAAAGCAAACTGGTCAACCTAATCAAACTTTTTCTAGCTTGTGGAAGGC
CTCTCAGAAAATATCTTTCAGAAGTAGCTCTGTGTGCTCAGACCCCTGGAACCAGTCATTCTGCTGCAACAGAGGCCTG
TGCTTTCCTATTGCCACCCATTTTCACTGCTCAGTGGCCTCTTCAGATATTAGTGCTATTTCCATAATTCTAGTTGCAT
ATGCTTTTTTGTTTCTTACCTATGCCACTTAAATATTGTGACATTTTGGGAGCACTTAAACATTTACTTCAAAACTTCA
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Fig. 6.10

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CATCTTTGTGGTATTTTTCTATAGTAACTTGTGGTTGTTAGAAATGCTGAAAATAAATGGTGCTGGGATAACTGGGTAG
TTAAATGTGGAACATTGAAACTGGACCCCTTTCTTTCACCATATAGCAAAATCAACTCAAGATGGATTAAGGATTTAAA
TATAAAACCTAAAACTATAAGAATCCTAGAAGAAACCCAGGAAATACCCTTCTGGACATCAGACCTGGCAAAGGTTTTA
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GGTCAAATATCCAGAATCTATAAGTAACTTAAACAAATCAACAAGCAAAAAAACAACCCATTAAAAGTAGGCAAAGGAC
ATGAACACTCATCAAAAGAAGAGAGATGAATGGATGAATGGCCAATAAGCATATAAAAATGCTCCACATCACTAATCAT
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ATAACATGCTGGTCAGGTTGCAGAGAAAAGGGAATGCTTATACAACTGCTGGTGGGAGGTAAATTAGTTCAGCCACAGT
GGAAAGCAGTTTGGCTATTTCTCAAAGAACTTAAAACAGAACTACCATTCAATCCAGCAATCCCATTACTGGGTATATG
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GAACAAGAGACACTGAGACCTAGTTGAGGGTGGAGGGTGGGAGGAGGGTGAGGATTGAAAACTACCTATGGAGTACTAT
GCTCATTTTCTGGGTGATGAAGTAATCTGTACCCCAAACTCCTATGACATGCAGTTTACCCATAAAACAAACCTGCACA
TGTACCCCCCGAACCTAAAATAAAGTTGGAATGGCAAAAAAGAAATGCTAAAAATATATTATTTCATTATCCCAATA
AGCTCTTACAGTAATTTTTTTATCTGTGTGGCTTTTTTGGGGGTATATCTCCAGTATAATTTAAAATAATCAGTTAAT
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GGGAGAAGGAGAAATTATGGGAATAGTGGTGGGATTAGGACTTTTATAACAAGAATGCAAAGAAGAACCAGTGTTCAGC
TTTTAATTACTGTAGGATGTTTCTAATAATGCTGGCAATGAGGAAAGCTTGACTTTGTCAGGAGAAGGAAGGATTTTCT
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Fig. 6.11

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Fig. 6.12

18/375

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Fig. 6.13

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TACTTCAGACCTCAGAATGGCAGATCCACTGACAGCTTGCTCTGTGTGCCTGGAAAAGCTGTAGGCACTCAACACCAGC
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TATAAGTTATAATTTTTCTTATTAATTATAAAGTATTCCCAGGGACTCTGTGAAGTCCTGTTATGTCTGGGCTCACGTA
TGAGGCACACAAGGCAGGCAGTAATGAGACTGCTTTTAATCTGTAAATATATAAGTAAACATTCAAATTAATTCCATTA
TTTTAAAGAGTGAGATTAAGAAATGTTAAAATGCACTCATTTTAACAATGTTTCCATTGTGCAAAACATTTTTGGCATT
ATTTGTCTTTTTCAGGCAGGAAAACCAGTGTCAAAACTTTATAATCATGTCTTATTTCTGTGTGTATGTGTCCACATAA
AGTATTAATTAGCATCATTGCTTGTTTCCAAATGCCTTTTGTCTGTTTCAAAACTCAAATCTACCATTAAAGGAAAAAA
AAATTCACACAATTGAGAATATTCCAAAGAATGGGCCTAATGGGGATCACAGAACTGATTGGAAGACTGAGACTGAAGA
TCTAGGCTCAGAAATTTGGCATGAATCTTGGCAGTAGGTGACACAGTTCAGGTCACACCACAAAGCAATCTGGTGTAGC
ATGTCTCACTGGCATGGCCACCATTAGACTTTATCCTCACTAGCACCAATTGGCAATGGAGTGTTGTACCACTTCCTTT
AGATGCAGTCATCCCAGGTGGGAGTGTTCGTGCAAGCCAAGCTTATGTGCCATGCACCAGATGTCAGTGGATGGGGGTG
TGTGAATCTGTTAGGTAGCAGAGCACAGTTTGCCACCAAAATATACACAATGTAGCATTTCCCCACAATTGCCCATTAA
AAATCTTTTACATTAAGCATTACTGACATTATAGAAGTGTCTTTGGATTCTTTTATTAGTTAAGGTGAGCATATATTAC
GTTTGTAATTTTAAGTCGAGTACATAATAATTATAAAAGAGAGATTTTTATTCTTTCAAGTTTCAGGTACAGAGATAGT
CTCTTTGCTTCGATATAGGGTACACTAGAGAGGGTTGTACACCTCAGTTCATTGCTCCACACTCTCCCAGCAGGGAGCA
TAGAACCTTGACCTTTAGAACATTCCACTTAATGGTGCAATAACCTGTATGCAGTGTAAGCTGGTAGGAGAAATCAAGT
AAGAAGAGGAAAGAAATGGCCATAAAATGCCCACACATTTCCTCTTCGCCCCCACCCGCCCCTCCAATCATAAGTGATG
```

Fig. 6.14

CTCCATTGGGCACTCTTTACCTCAACTTAAGTTTCACCTTTTTCTGGTTTGGAAAGGGAAAACTACGTTTTTTTGTTTT
GAGACGGAGTCTTGCTCTGTCACCCAGGCTGGAGTGCACTGGTGCGATCTCGGCTCACTGCATCCTCCCTCCCAGGTTC
AAGTGATTCTCCTGCCTCAGCCTCCTGAGTAGCTGGGATTACAGGTGCCCGCCACCACACCCGGTTAATTTTTGTATTT
TTAGTAGAGACGGGATTTCACCATGTTGGCCAGGATGGTCACGAACTCCTGACCTAAGGTGATCTGCCCTCCTTGGCCT
CCCAAAGTACTGGGATTACAGGCGTGAGCCACCGCGCCCATCCAAACTACATTTTTAAAAAAAGTTAAAAAGCAAAAAC
GAAACAAAAGTTGGGAAGGCAAGTGTTCTGTTCCATTTCATTAGGAGGCTGACTCTCTAGGAACCTCTCCTGGTCGCTG
GGACCATTGTTACAAGGTTGAGAACTGTGAGTGTACGGAAACCCAGGGGAAACATTTTTCCTCATTTACTTCTCAAAAT
AGAGGTAGGAATAAGGCATGGAAAAATATAAATGAGTAAATATAATCCTTGTTACCTTGGAATAATTGGTGATAAAGTG
CAGGTGCATCTTATTTTCATTCTATCAAGTCGTTTTCTAATTGGATGTTCTTGAAACAGTGGCTCATTGCTTGTACTTG
TTCACTCCTATTTTCTTGCTACTATCCTGTTTTGGGGGCAGGGCAACTCTACCCTGGGGAACGATCAAATGTTTAGGCA
AATATATCCTGGTGTTATAAGCTCATTAAATTTCTTGTCATTTCTTGCAGTGTGCTGGGAAGTGCTGTTTGTGGCCCGT
TTTTGCAATGTAGAACCTCTCTTTTGTGAGAGCCATGGAAGTTCCTTATTTGATAATTAGTCTCTTGCTTTAGGCAGTG
CCATTAAGAACTCCATATAAGCTGCCAACACTTTGTGGATGTGCAGAACCCCGTAAGCTTCCTATAGCTACTATTGCAA
GTAAAATTTTCAGATGCTTCTGAAGTCTTACCCACTTATCTTGGCGTAGAGGTTCTCAAGCTTCAGCACTGGAGGATCA
CCTGGAATTGCTCGTTAAGACACACATTGCTGGGATCCACCTCCAGGGTGTCAGAGTCAAGTCTGGGGTGAAGCCTGAG
AATGTGCATGTCTAACCAGTTCCCAGGTGATGCTGATGTTGCTGGTTTTGAGACTATATTTTGAGATTAAGAACTGCCT
TGGATTAAGAGCTGACCTGCGATCTACAGCTCAAATAGTGAAGTAAACATCCTAAAGAAAATGGAAAAACCAGTGCAGT
GAGTGATTGAATTACTATTTGTTCAATATCACAGAGAGCATAGTATTACATAAGGGCTTTGGGGAATATTTTAGGTAAG
TATATATAACTCTTGCCACCTAGGATGTTCATAGTAACTATAAGACAGTATTTTTGTTTTCCAAGTAATTTTAATGATC
CTGTAGATCCTCTTCCTTTTTGTATATAATATCAATCTAATAGTTTCTTGTTTAATATAAAAATGAAATCTTATTTTAC
ACAAGCAGCAAGCAGCTATTTTTTCAGATTTTCCCCCTATAATCTAAGGGAAAGTTATTTTAAAATAGAAAAGATGTGG
GCTTCAAAAAAAGCTTTGCAAATATGTTGCAATAATACGAATGATTTCAGTGTTGAAATCCATTTGTGAAAGCAGGCTT
TGCTTATATTTTGGGTCCTGCCTTCTATAAAATGCTCAGATTTGCTTTTATTAAGATCATACACTCAGTGACCTGAGGA
CCAGATGGAGGTTATAAGCAGCTCTTTAAGGCTTCAGAGCTTAGCCTAGAGAGTCAAACAGCTCTTTGAACTGGCGTCT
CAGCTCTGTCATTTAGGCAAATTGTTTATTTTCTGCAAAGAACCAGCCTTAGGCTTTGTTGATCTACGTATGTATATTT
GTTTTTTATTTATTTGCATTATTTCATTCATTCTGTTTTTCTTTGACTTTAATTTGCTAGTTTTCTTTGAAGCTTGTGAG
ATAAATGCTTAGATCATTGATTATCAAAATTGTGTCTATTCTAATATATATATAATATATATATGATAAGGCTATACAT
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TATTCTAATTTTATTATGATTTCTTGAGCCTATGGATTATATAGAAATATATTTCTTAAATTGTAAACATATGGGGATT
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ACATGTGCACAATGTGCAGGTTTGTTACATATGTATACATGTGCCATGATGGTGTGCTGCACCCATTAACTCGTCATTT
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TCCTGTGTCCATGTGTTCTCATTGTTCAGTTCCCACCTATGAGTGAGAACATGCGGTGTTTGGTTTTTTGTCCTTGCAA
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TAGTATTCCATGGTGTATAGTATTCCATGGTGTATATGTGCCACATTTTCTTAATCCAGTCTATCACTTTTGGACATTT
GGCTTGGTTCGAAGTCTTTGCTATTGTGAATAGTGCTGCAATAAACATATGTGTGCATGTGTCTTTATAGCAGCATGAT
TTATAATCCTTTGGGTATATACCCAGTAATGGGATGGCTGGGTCAAATGGCATTTCTAGTTCTAGATCCTTAAGGAATC
GCCACACTGATTTCCACAATGGTTGAACTAGTTTACAGTCCCACCAACAGTGTGAAAGTGTTTCTATTTCTCCACATCC
TCTCCAGCACCTGTTGTTTCCTGACTTTTTAATGATTGCCATTCTAACTGGTGTGAGATGGTATCTCATTGTGGTTTTG
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GATTGTGGATATTAGCCCTTTGTCAGATGAGTGGGTTGCAAAAATTTTCTCCCATTCAGTAGGTTGCCTGTTGACTCTG
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TTTTAGGTCTAGCATTTAAGTCTTTAATCCATCTTGAATTAATTTATGTGTAAGGTGTAAGGAAGGCATCCAGTTTCAG
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GGTTTGTCAAAGATCAGAAAGTTGTAGATATGCAGCATTATTTCTGAGGGCTCTGTTCTGTTCCATTGGTCTATATCTC
TGTTTTGGTACCAATACCATGCTATTTTGGTTACTGTAGCCTTGTAGTATAGTTTGAAGTCAGGTAGCATGATGCCTCC
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TCCATTTCTGTGAGGAAAGTCATTGGTAGCTCGATGGGGATGGCATTGAATCTATAAATTACCTTCGGCAGTGTGGCCG
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GCAATTGTGAATGGGAGTTCACTGATGATTTGGCTCTCCGTTTGTTATTGGTGTATAAGAATGCCTGTGATTTTTGCAC
ATTGATTTTGTATCCTGAGACTTTTCTGAAGTTGTTTATGAGCTTAAGGAGATTTTGGTCTGAGACGATGGGGTTTTCT
AGATATACAATCATGTCATCTGCTAACAGGGACAATTTGACTTCCTCTTTTCCTAATTGAATGCCCTTTATTTCCTTCT
CCTGCCTGATTGCCATGGCTAGAACTTCCAACACTATGTTGAATAGGAGTGGTGAGAGAGGGCATCCCTCTGTCTTGTG
CCAGTTTTCAAAGGGAATGCTTCCAGTTTTTGTCCATTCGGTATGATATTGGCTGTGGGTTTGTCATACATAGCTCTTA
TTATTTTGAGATACGTCCCATCAATACCTAATTTATTGAGAGTTTTTAGCATGAAGGGTTGTTGAATTCTGTCAAAGGC
CTTTTCTGCATCTATTGAGATAATCATGTGGTTTTTGTCTTTGATTCTGTTTATATGCTGGATTACGTTTATTGATTTT
CATATGTTGAACCAGCCTTGCATCCCAGGGATGAAGCCCACTTGATCATGGTGGATAAGCTTTTTGATGTGCTGCTTGA

Fig. 6.15

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TTCGGTTTGCCAGCATTTTATTGAGGATTTTTGCATCAGTGTTCATCAAGGATATTGGTGTAAAATTCTCTTTTTTTGT
TGTGTCTCTGCCAGGCTTTGGTATCAGGATGATGCTGGCCTCATAAAATGAGTTAGGGAGGATGCCCTCTTTTTCTATT
GATTGGAATATTTTCAGAAGGAATGGTACCAGCTCCTCCTTGTACCTCTGGTAGAATTCGGCTGTGAATGCGTCTGGTC
CTGGACTTTTTTTGGTTGGTAAGCTATTATTATTGCCTCAATATCAGAGTCTGTTTTTGGTCTTTTCAGAGATTCAACT
TCTTCCTGATTTAGTCTTGGGAGGGTGTATGTGTCCAGGAATTTATCCATTTTTTTCTAGATTTTCTAGTTTATTTGTG
TAGAGGTGTTTATAGTATTCTCTGATGGTAGTTTGTATTTATGTGGGATCGGTGGTGATATCCCCTTTGTCATTTTTTA
TTGCATCTATTTGATTCTTCTCTCTTTTCTTCTTTATTAGTCTTGCTAGCGGTCTATCAATTTGTTGATCTTTTCAAAA
AACCAGCCTCTGGATTCATTGATTTTTTGAAGGGTTTTTTGTGTCGCTATTTCCTTCAGTTCTGCTCTGATCTTAGTTA
TTTCTTGCCTTCTGCTAGCTTTTGAATGTGTTTGCTCTTGCTTCTCTAGTTCTTTTAATTGTGATGTTAGGGTGTCAAT
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AGATTCTGGTATGTTGTGTCTTTGTTCTTATTGGTTTCAAAGAACATCTTTATTTCTGCCTTCATTTTGTTATGTACTC
AGTAGTCATTCAGGAGCAGGTTGTTCAGTTTCCATGTAGTTGAGCGGTTTTGGGTGAGTTTCTTAATCCTGAGTTCTAG
TTTGATTGCACTGCGGTCTGAGAGACAGTTTGTCATAATTTCTGTTCTTTTACATTTGCTGAGGAGAGCTTTACTTCCA
ACTATGTGGTCAATTTTGGAATAGGTGTGATGTGGTGCTGAGAAGAATGTATATTCTTTTGATTGGGGTGGAGAGTTC
TGTAGATGTCTATTAGGTCTGCTTGGTGCAGAGCTGAGTTCACTTCCTGGGTATCCTTGTTAACTTTCTGTCTCATGGA
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GGTTGCAACCTGTGACTGTTTTTGTTTTCCATGTGCTTGGTAGATCTTCCTCCATCCCTTATTTTGAGCCTATGTGTGT
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GCTGGTACCGGTTGTTCCTTTCCATGTTTAGTGCTTCCTTCAGGAGCTCTTTTAGGGCAGGACTGGTGGTGACAAAATC
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GGTTGAAAATCCTTTTCTTTAAGAATGTTCAATATTGGCCCCCACTCTCTTCTGGGTTGTAGAGTTTCTGCAGAGATAT
CCGCTATTAGTCTGATGGGCTTCCCTTTGTGGGTAACCCGATGTTTGTCTCTGGCTACCCTTAACATTTTTTCCTTCAT
TTCAACTTTGGTGAATCTGACAACTATGTGTCTTGGAGTTGCTCTTCTCGAGGAGTATCTTTGTGGCATTCTCTGTATT
TCCTGAATTTGAATGTTGGCCTGCCTTGCTAGATTGGGGAAGTTCTCCTGGATAATATTCTGCAGAGTGCTTTCCAACT
TGGTTCCATTCTCCCCGTCACTTTCAGGTACACCAATGAGACGTAGATTTGGTCTTTTCACATAGTCTCATATTTCTTG
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TACCCTTTCTTCCAGTTGATCGAATCGGCTACTGAAGCTTGTGGATGCATCACTTAGTTCTCGTGCCATGGTTTTCAGC
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GGTGGGGCGCCCCTCCCCCAACCTTGCTGCTGCCTTGCAGTTCGATCTCAGACTGCTGTGCTAGCAATGAGCGAGGCTC
CGTGGGCGTGGGACTCTCCGAGCCAGGCNCGGGATATAATCTCCTGATATGCCGTTTGCTAAGACCATTGGAAAAGTGC
AGTATTAGAGTCGGAGTGACCTGATTTTCCAGGTGCCGTCTGTCACAGCTTCGCTTGGCTAGGAAAGGGAATTCCCTGA
CCCCTTGTGCTTCGCAGGTGAGGCGATGCCTCACCCTGCTTCAGCTCACACTCGGTGCACTGCACCCACTGTCCTGCAC
CCTGCAGTGTAGCAAAGTTTGAAATGTAGATAGCCAGAAGTCAGTCTGGGGGAAAGCAATTCCAATTTTCAAGCTTGTA
AAATTGTTTTCATTGATTTCTACTCAAATTTGAAATTCTTTCCTGTCGTCAACATGTTTGATGATGCCTAATATGAGCT
ACAAATTCACTATACCATTTTTTTCTTTTTTGCTGGAACATGTATAAAATAGAATATATCTGAATTTCCATGTAGGGCT
GAAACATGCTGTCATAATAGAAACAATATGCCTACATCTCCGTCTCTCTTTCTGTGTGTGTGTGTATGTATGTGTGAAG
TCATATGTTTTATGCTTTCCTATATATCAGATTATGTTTTAGCATTTCAGAGGCACTGTGTCCTGCTAAAATCCTGTGT
TTCCAGATGAAATGGCAAACATTATTTCCAGCAATGTGATAAAACAACCAAAGAAAGTGTTCAAGTGACCCGATTCATA
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TAGCTTAACACAGTAAAGTATCACTTTTAAATGCATTTGCATTTCTCATGGGTGAGGGTTAGCAAATTCTGACCTGGAA
ACCAAATCCAGCCTACTGTTTATTTGTAAAAAATTACATTTGAACACAGTCCTACCCATTTATTAACTTACTGTTTAT
CACTGCTTTTGTTCTAGAAAAGCAGAGTTGAGTAGTTGAAAAAAAGACTGTGGCCAGGCACAATGTCTCATACCATAAT
CCCAGCACTTTGGGAGGCCAAGGTGGATCACTAACTGGTCACTAACTCTTGGGCTCAAGTGGGTGGATTACTTTAGCCC
AGGTGTTAGTGACCAGCCTGGGCCACATGGTGAAACCCTGTTTCTACAAAAAGTACAAAAATTAGCCGGGAGTGGTGGT
GTGAACCTGTAGTCCCAGCTACTGGGGAGGCTGAGGTGGGAGGATGACCTGAGCCTGGGAGGTCAAGTAAGGCTGCAGT
GAACTGAGATCATGCCACTGCACTCCAGCCTGAGTGACAAAGTGAGATCCTATCTCCAAAAAAAAGAAAAAAAAAAAGA
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Fig. 6.16
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CTGTAATATAAGACCCCCAATGTCAAAAATATTTACTATCTGATCCTTTACAAAAGCGTTTGCTAACTCCTGTCCTTGT
GTATGCTTTGATTTCATATTAATTATATCTTATGTTATTTATGATACATATATGANATACACATATATATCTAATATGT
TATATATTTAAGAATTATTCACTATGCCACAAACTGTAGATACAAAAGAAGTATCTCTAGGGAAAGCCAAAAACAAACA
TGGAAAGAAAGAAGGCTAGCAGAAGCTTCAAAATATCAAAAACTCTCTTACTGTGTGGCAATATAAACTAAAAACTGAT
GCTCAAAATCATGAAGATAGGAAAAAGAATCAAGACAACCAAAAAATATGGTTAATAAAAATGAAGACAGAAGACATCA
AGATGTTTATGAGAACCAGGGTCCTTTCAAACAGGGGGTGAAAAAGGAGTGAGAAGATGTGCATGGCAGTGAAGAAGAA
CTTAATTTGCACACAGTCAACGGGGGGGTGGGCAGGAGGGCCAACACAGCTGCATTTCTACCACATTATTTATTTTGGT
AAGTACTGTCATCTTTTGTTATGAGCCATATTTCATTCAACAGTGTTTAATGACTTTTTTCCAAATCATTACTCCATAA
TATTTACTGAGAATCAAACAGAACTTACAATAGAAAAAGAGATACTTCTTAATTAGATATTTTGGAAAAAAATCATTG
TCGTGTGACAAGACAACCAAAGAGTAGTCAGCTAAACCTATAGAAATGAAGTATTCTAAAAATGTGTCAGACAGCTAAT
TAACACAATATTTTATTTTTCTAAAGATCTTGCAGTGTTTGTCATCTTTATCACCTTTTAAAAGTTTGCATTTTATTGT
CACTTTTAAAATTTTTAAATAAATGTCCATTTTGTATCTATCATTCTGTGTCTTTATGTAATAGTATCTAATATTCTAT
ATCTGTATCAAATATTCTGTATGTACTTCATATTCATTTTATCTTATTTTATTTATTTATTTATTTATTTATTTA
TTTATTTTAGACAGAGTCTTACTCTGTCTTCCAGGCTGGAGTGCAGTAGCACAATCTTGGCTCACCACAACCTCCACCT
CCCGGGTTCAAGCAATTACCCTGCCTCAGCCTCCTGAGTAGCTGGGGTTACAGGCACACGCTACCATGTCCAGGTAATT
TTTGTATTTTTAGTAGAGATGGGGTTTTGCCATGTTGACCAGGCTGGTCTCAAACTCCTGACCTCAAGTGATCTGCCCCC
CTTGGCTTCCCAAAGTGCTGGAATTACAGATATGAGCCACCACATCCAGCCCCATTTTACCTTAATTTGTAAGAATATA
TAATTATATTTATAATTTTATTCATTTTATATTAATTTTTCTATTTCTTAAAGGCTAAATCTCTAGTTTCCTTTTTATT
TTTCAGGTAGAGGATTTTAATTCACTGAGTACCTGATCATTTTACATACCTAGTCAAATAAATATGAAATTTGAAGAAC
TAAGATTAAGTTTTAAAAATTCTTAAGTGCATACAATCCATTGAAGAAAACAAAAATTGCTTAAGCAGTTTTTAAAACC
TTGATAAGAGGTCATCAACATGATTTAAATGTGAAATTAGCTAATGTAAGTAGCAGTTAGAATGAGATTTAGCTTATAT
ATTTGGAGGTATAATGGAAAACTCTGGTGAGGACTCTTTTATTGCACACCAAGAGAAACATGAGAAAGAAAAATGAGTT
GCATTTTTCTGCATTTTTCCAGAAGTGAAATGCTTTGCAACATCCTCAGAAAGAAAGAGAATACATGAGTAATATGAAA
GAAAACTATATAAAGGATTTAATTATTTCCAAATTGCCTGGGTGTTTTTTTTAGTGCCTGCTTTTTTTTTCAAGCAGGAA
TAACTGTAGAAACTTTTATTTATTTATTTTTAAAGGAATGGTTACTATTCTTAGGAAAAGTGGCAGTAAATATAGTTAA
TAACGGTAATTATAATTTTATAAACTCATTCAAAAGTCTTGATTTTAAAGGCGATAGTAAAAAAAATATATATCTATTT
ATCTTGTTCCCTGAAAAATGGTAGCACTCCTGACCACTGAGAGACTGTTCTCCTCGGCATAATTGATGGCCTTCAAAGC
CATCATACTTCAGTATTCAATGTCAAAGATTTATTTCCATTTGCAAAATTTGCTTAGAACTCACCTATATTTACCTTTC
CTCTTATGACTTATAAGATAATATTAACATCTACATAACAGAATCTCACTCACAGAATGTTTGAGCCACGAAGGTGCTC
TGAGACCCTCTGCCTCAACAGCCTCATTCATAGATGAGAGAAGCCCAGAGAGGTGAGGACGCCTTCCAAAGCTCTAGCA
CTTGTTTCTAGACATGGAACCAGGACAGCCCTTGGCATTTCCCACCACATTTTCCTGCTTATTATGTGGCATGAGCTTG
TCTTTAACACTGAATTAGAATTATTGCACATTACATTGCATTCATCAAGGCCTACCCTTCAGGCAGTCTGATGTAACAG
TCTATTATAAGACATCCGAGAAAAACCATCCACTATCCTTTCAGATGTTCTCAAAACAATTTGTATTAAGAGCTAATAA
AATGAAACAGAAATGCCAGGTAAACTTTTTAAAATAAAGATTTTGTGATTCATTGGGTTTAATTATTGGTTTTATTTT
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ATTATGATATAATTTACATACATTTAGATATATTTAAAGTGTACAGTTCATTTATTTAGTATAACCACAGAGTTGTGCA
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CAGCTCCCTCTTGCCACCCTGTACCCTCAGTCCTAGGCAACCACTAAACAACTTTCTGTCCTCATAGACTTGTCTATTC
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TTTCACATTTATCATGTAGCATTTGTAAGTGTCTTTCCCTTTTATAGCTGAGTAATATTCCATTGTATGTGTATTCCAC
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TTGTGGAAAAAATTAGAACTTTTCTAAAGGACATTATTTTTAAAGTTTATTACTTATTATTTATTAATTTTTATTAATA
CATAATATTGCTATGTATCAGGTACATGTGATAGTTTGATACATGCATACAGTATATAATGATCAAATCAGCATATTTA
GAAAATCCATTACCTCAAGCATTTATTTCTTTGTGTTTGGAACATTTCAGAACTTCTCTTTCAGCTATTTTGAAATATG
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TTGTACCCATTAACCAATCTGTCCTCATCCCCCCAGCCCTTTCCCAGCCTCTGGTATCTATCATTCTATTCCCTACCTC
CATGAGGTCAACTTTTTAGCTCCCACATATGAGTGGGAACATGTGATACTTGTTTTTCTGTGCCTGGTTTATTTCACTA
AACATAATGACTTGCAATTCCATCCCTGTTGCCGCATATGATGAGATTTCATTTTTAATGGCTGAATAGTATTTTGTTG
TGTATATATACCATATTTTCTTTATTCATTCATCTGTTGATGGACACTTAGGTTGATTTCATACCTTGGCTGTTGTGAA
TAGAGCAGCAATAAATATGGGGGTACAGTTGTCCCTTTGATTTATTGATTTCCTTTCCTTTGGACAGAGAGACAGTAGT
GGGATTGCTGGGTCAGATGGTAGTTCTATTTTTAGTTTTTAAAGACACTTTCATACTGTTTTCCATAGTCGTTGTACTA
ATCTACATTCCCAACAATGCATAAAGAGTTCCCTTTTCTCCACATAATCACCAGCATGTGTTATTTTTTGACTTTGATA
ATAACCATTCTAACTGGGGTGAGATGGTATCTCATTGTGGTTTTGATCTGTATTTTCCTGATGATCCGTGATGTTGAGC
AGTTTTTCCTAAACCTGTTAGCCATTTGTCTTTTTAGAAATGTCTATTCATGTCCTTTGCTCACTTTTTAGTGAGATTA
TTTGATTCTTTGCTGTCGAATTGTTTTGAGTTCTATGTATATTCTGGATATTAGTCCCTTGTTGGATGAATAGTTAGCA
ATTTTTTCTCCATTGTTCATGTTATCTCTTCACTCTGTTGATTGTTTCCTTTCCTGCGCAGAAGCTTTTTAGTTTAATG
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Fig. 6.17

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TCGTTCCATTTGTCTATTTTTGTTTTGGTTGCCCCTGCTTTTGAGATCTTAGCCATAAAATCTTTGCCTAGATCAATAT
CTTGAAGCATTTCCTCTATGTTTTCTTTTAGTAGTTTTATAGTTTCAGGTCTTGTATTTAA CTTTAATCCATTTTGA
ATTGATTTTTATACATTGTGAGAGATAGAGGTCTAGTTTCATTCTTCTGCATGTGAATATCCAGTTTTCTTAGCACAAT
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TTATTTCTGGATTCTCTGTTCTGTTCCACTGGTCTGTGTATCTCGTTTTTATACCAATAGCATAGTGTTTTGGTTGCTAT
AGCTTTGTAGTATATTCTGAAATGTGTTAGTGTGATTGCCTTCAGCTTTGTTCTTTTTGCTGAGTATTGCTGCTATTTG
GGCTCTTCTATGGTTCTATGTGAATTCTAGGATTGTTTTTTCTATTGATTCAAAGAATGTCATTAGTATCTTGATAGGA
ATTGCATTAAATCTATAGCTTACCTTGGGTAGTATAGTCATTTTTAACAATATTAATTGTTCCAATTCATGAGCATAAT
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CATAGAAATCTTTCACCTCCTTGGCTAAATTTATCCCTGAGACTTTTTGAAGTTATAAATGAGGTTGCTTTCTTGGTT
TCTTTTTCAGATAGTTGGTTATTGGTGTATAAAAACACAACCGATTTTTTATATTGATTTTGTGTCTTGTAACATTACT
GAATTTGTAAAATCATGTCATCTGAAGTAGAGACAATTTGGCTTCTTCCTTTTTTTTTTTTTTTGAATTGGAATCTCTCT
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GCCTCCCAAAGGGCTGTGATTACAGGCATGAGACACCGCATCTGACCCCTTTACTATACCTTGTAGTGTCTCCCCTACA
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```

Fig. 6.18

24/375

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Fig. 6.19

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Fig. 6.20

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TGAAAAGATCCCAGGAGCTTCCCAATACATATGCAATCTCTGTAAAAGCAGTGTGTTTGCAAACACAGCTGCAAGAAAA
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TGTCTGGAGTTTGTTCCTTCTGATGTTTGGATGTGTTCGGAGTTTCTTTCTTCTGGTGGGTTCACGGTCTCGCTGGCTC
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GGGTTCGCGGTTTTGCTAGCTTCAGGGGTGAAGCTGCAGACCTTCATNGTGAGTGTTACAGCTCATAAAGGCAGTGTGG
ACCCAAAGAGTGAGCAACAACAAGATTTATTGCAAGAGTGAAAGAACAAAGCTACCACAGCATGAAAGGGAACCTCAG
CAGGTTCCCACTGCTGGCTTGGGCAGCCTGCTTTTATTCTCTTATCTGGCCCCACCCACGTCCTGCTGATTGGTCCATT
TTAAAGAGAGCCGATTGGTCTGTTTTACAGAGAGCTGATTGGTCTGTTTTGACAGGGTGCTGATTGGTGCGTTTACAAT
CCCCGAGCTAGACACAAAAGTTCTCTACCTCCCCACCAGATTAGCTAGATAACAGCATCCATTGGTGTATTTACAAACC
CTGAGCTAGACACAGAGTGCTGATTGGTATGTTTACAAACCTTGAGCTAGATACAGAGTGCTGATTGGTGTATTTACAA
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Fig. 6.21

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TCCCTTAGCTAGACATAAAGATTCTCCAAGTACCCACCAGACTCAGGAGCCCAGCTGGCTTCACCCAGTGGATCCCGCA
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GCTCTTCAACATCCCATAACCTATGCATTTAGTTGCTGTTCATAAATAGTCACCTGCATATGTTCATCTTGCTTTTACC
TGTGTTCATCCCATCTTTTTTGTGAAACTCCTGGAAGGCAGGTCGTGTCTTCTTTCATTCTTTTATACCATCTCAGAGG
GCTGTAACAATACATGCTCAGAAAACATTTCTGTTCAACCTGCATGAAGTACAAGGCAGGCTGAAATGTAAAGAAATAC
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CCACTGGGACCTTGCTTCCTACGAGGCTTGTCCTGCATTTCATGGCCTTAAATTTCACCAATGTAGTTATCTAGTGGTG
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Fig. 6.22

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CTGCCTACATAATTTCCTTAGAATTAAAGCATCAATAATGGACCTTTGCAAAAATAAGCACTTTCTAACACCAATGTGC
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Fig. 6.25

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GTGGCATAGCTCAGGGTAAAGTGGCTCTCCTCACCTCTCAATACCCTACTGTCTTTTGCCTAAATTCCAGTTAAGGATG
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Fig. 6.25
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31/375

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Fig. 6.26

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Fig. 6.(27.)

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TGTTCCTTCTTGGAACATATGTTATGGAACTTGACTAATCGCCCTGGCAATATGTTGCCAAAGTTTTCCTGGCAATAGA
TCAAAGTCTGTAATATACCAGCACATCAGCTTTGCAGAGCTCAGACAATGAGAAAACAGGTAGGGGTGGGAGAAAGCAC
TTGCAATTAATTTTGGAGTCAGATTTTGATAGAAAAGTAGAAAGAGGATAGGACTAATTGACTTCTGAAAGAATTACAN
GTGGCTGGTGTGCCCCATGACTCACACTGTAATCTAAATACTTTGGGAGGCCAAGGTGGGAGGGTCTCATGAGGCCAGG
```

Fig. 6.28

GGTTTGAGATCAGCCTGGGGAACATAGTGAGACCCTGTCTCTACAAAAAATAAAAAAAATTAGCCAGGCTTAGTGGCAT
GTGCCTATGATCCCAGCTGCTTGGGAGGTTGAGGTGGGAGGATCACTTGCAACTGGGAGGTTGAGGCTGCAGTGAACCG
TGTTCACACCACTGCACTCCAGCCTGGACAACAGAGCAAGATCCTGCCTCAAAAACAAAAACAAAAACAAAAAAAAAGA
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TAAAGATAACAATGCTATCATTTCCCCACATCATTTTGGCAGAAATGTGAATGGATGGTTAAACTCGGTGTGCCTGTGA
GTTTGATGAAATGGATGTGCTCATACAACTACCTGATACAACCTTTTGGAGGGCAGGTGGCTGATACTTACTGAAATGT
GAAATGTATAATCTTCTTGCCCAGAAATTCTGGGTCTAGGACTTATCCTAAGAAGACAACTATACAAATGTAAAAATAC
ATTTATGTAGTATGTTTGCAAAAGCATATTCATGGGGAAAATGAGAAATAACTTTAATGTTTATCAATAGAGAACTGGT
ATAGTAAATTATGATAAATACCTACTGTGAAATCCTGCATAGTCAATAAAATGATGGTGTAAAGCTTCATATATTAATG
TGAAAAATTATTTAATGGTACAAAACAGGTTATGAAACACAACAATCACATTTCTATACAGTCATATTACTAGAAATAT
GGGAAGTTGTTAATTTAAATCTGAACAGTTAGAATTTCAGGCTACCTTTACTTCCTTTGCTATTTAACTGTTTCCTCTC
CCACCTATCAACTCTATGTCCTTAGGGAAACTTATTTAACCTTTCTCACTCTTGGTGAAAATGGAGATAATACCAAACT
CTTAGAGGTTGTTGTGAGAATCAAGCAATAAAATACATGTCAAGCACCTAGTAAAAAATGTATTTTGATTTTTTTTTGT
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CTAAGTGGGCTTTTGAATAACTTCCACTTATTTTAATGCAGGGAAAGCTCTAGTTCCAAATATGTGGAACAGTATTTTC
CTAAGGCTGCAATGTGTCAGAAAAAGACCATATTGAGTCATTGTTTTCTGAAGCCAAGACAGGAGATTGAGAGACGGAT
GCTGATGGGCAGCCTAAAATAATATTTCATTCTTGTCATCCTTAGGCAGTTTAAGACCAAGTCATTTTGGATAGATCTA
CAGTTTCCCTCTCGGTTTGGAAGAACAAAGCTTTTGTTATTTATGTTCTTAAACAACAGCGCTTTGTCCTATACTCAGG
AGCGTGCAAAGCAAACAGAAGATTTCCTGAACGGCCATCCCATTTATGGTGAGGAGTCAGTCCAAGAGGTAAGTAGACA
TACTCAGCTGCAACCATCTGCAGAGGGCCCACATTGGCAGCAAAAGGAAATAGCCAAGACTCACAGGAGCACAACAGCC
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AATACAGGGCAGGGTTAAAGTGTCAATTGAGAACCAGGCAAAAAACTGATGATTAATAGTGACCAACACACTTCACTG
TGCTGTTATCTTAGAGGCCTTAAGGCAAAATCTTACTTACAAATATTAGTTGAAATGCAATATGTGATGTTAGTAGACA
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Fig. 6.29 7

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ATACTTTAAGTACTAGGGTACATGTGCACAATGTGCAGATTTGTTACATAGGCATACATGTGCCATGTTGGTTTGCTGC
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TGCATTATTCTACTAAAAATAAAAAGAAAGATGAAAATTACCTCAGGCGTTTGCTGCCGTGCCTGCCCTTTGGTTTCTG
GGACGGCTCGGGTCCCGTAGCGCCGGCACAGCTGAGATTGCCAAGCCGGGAAGAGACCTTGCTCCAGGTGTAGCTGCGT
TTTCCCCAGATCACCTGTCCTTTTCCCCTCCGACAAGGAAGCTGTGATTTTTCTCTGGCCTTTAGAGGCAAAGTGATTC
CAGATAAGTAGATTAATGTGTAGAATATCTCATCTGTGTTGTTCCAGTGCAGCCCTTTCAGCTTTCCAGAGCCAGTTAG
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Fig. 6.30

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ACTTGTTATGAGGAGCTAAGTGATTGGCTGGCTCTGGAGCTCAGTTTCATAGATTATAGCCCAGCGTACGAGAAGCACG
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GCAGCCAATTGAAGACATACACTATGAATAACTAAAACATTTTTATATGAACAAAAATGCTCTTCAGTGGTTCTGTTTA
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CAACCATAACAGAAGAAAGTATGTGGATTGGAATTTGTGATTGGAGCAGATCAAATTAAGCCAGGGAAGCCGTTATTAG
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ATTTTCTATGTTAATAATGAAGGAATGACCAATCTGTAGTATATGCAAAAAGTACTGGGTAGAAATATATTAATTTTCT
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TGTTGGGTGTTAAGCCAATAATAAACTCAGCTATTTTAACAACCAAATTATTTTCCTCCTATAAACCAAATTTACAATC
AAGCTACTTAAGAAATAAAGACAACAAATAGAAACTGTTTTAATCTTCATATCCTGACCTAAAAAAATAGAAGCACGAAC
TCATGTCACATTCTCATATTTATCTGATTAGACTAGGAATGCCTAAAAAGAATTTTCCCATAACTCCATTTACTTGTTG
TTTATTTCCCTAGCTCATTCCAAAAGCAATTTAAAATACAACACCAAAAGAGTAAAATATCCTCAATCTGACATCCTAG
CAATCTTTTTATTTTGGTGAAACTGAACAATAATAATAGGCACTTGCATTCTTCACCCTGAGGCACTGAAAACACTGAT
ATGTCTCAGTTTTAAGATTCTGAAGATTATATTCTATTTAATAGATATTTTAAGTATATGTATTATATTTCTAGTCTTTT
GAACTAAGGCAGTCCACATATAGGTATAAAGTAATGCCTCAGCATAGTGTAGCAATTATTGGTGCAAAAGGTTGGTGCA
AAAGTAATTGTGGTTTTTGTCATGAAAATGGCAAGAACTGCAATTACTTTTGCATCAACCCATAACTCTGTGGTCTCTG
GAGCCAGACTGCCTGGGTTCAGATCCCAGCTCTGCCACTTAGTACCTGAGTTACTCTGGACAAACTAATTAATCTTCAT
GTACCTCCATTTAATTATCTGGAAAATGGGATAACAATAGTACCTATCTGGTTGGGTTGTTGTAGAGATTAAGTAAATT
AATATACGTAAAATGCTTTACATTTTACGTATATTAATTTCAAAATTGCTTCGAATGATGCATATCTCATAGTGCTTTA
TAAATGTTTGTCATCTTATTATTATTTGGGGAACTGATTATTTCACAAAATATATTTTCTAGATAGCTGAAGTTTATTC
AGTCAGTTTCAAAACTTCTAGATATAGCCAGTTTTCATGGGTCTTTACAAAAGGATCCCTGTATCTTTGTTTCTTCTGT
TTTAGAGAGATTAAATGGAATCTTCCTTTGGTGATACTGATTGAGGTGATACTGAGATAATTAAGAGGACATGAAGGGA
TTTAGGAAGAGTGTTTGCCCCTGCACTAACTGGCCCAGACTGCTCTACTGCATCAGCCCTCTGGCGATGTTCTAAGGGT
TTGGACTTCAGTCTACTGCCTATACTAGTACCCTTGCTTTAAAATTCCTGCTCCTATTTTGCTATAGGCTGTAAGCAAG
AAAATAGCCTAGTTTCTAAAACCTATGTATAAAATGAGAGAATAGGTGTCACTCAGGGTGTCAGTTCTCCCCTGAGAGT
TAGGCCAAATCTGTTTTTTGTAAATAAAGTTTTATTGTAACACAGCCATGCTCATTTGTTTATATGTCATCTACTGCTG
CTTTCACACACTGTAAGGGCAGAATTGAGTGGTTCCCGCAGAGACCATATGGTCTGTATAGCCTAAAGTATTTAATATC
TGACCTTTTACAAAAGAGTTTACCAAACTCTGTTCTAGAGCTTTAAAATTGGACAGAGACAGAAATAAATGAGATGTTA
CTACATTTAGTAGTCTCCACTAACCTGCCTATTCAATTATTTGCAGCACCACATCCCTTAAATGTATTTTGGAGAAACA
AGAATTTATTGCAGTAGGCTATGTTGGATAAAGAACAAACAGCCCTCAGTAGCACATTGTGAATATTAAGAGGTGGTGA
AGCTGTGATTGTCAAGCCAAGTTGTTTGGGAATCTTCATTTTAGTATTTGTTGCTCTAGTGTGATCAGTANATAACATG
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ATGCAATTATTCAAAGTAAATGTAAATATGTCTGTTGTCCTTTTCTTTTACATGAAAACTTCCCCCCTCCCCAACGCTG
TAAGATTCAGAAGCATAGTGTACAGAAAAGGATCTTGAACTTGGGTTTGAATCCTGACTTAGTCACTTTATAAAGGAGA
AACATCAAGTTACATTGCCACCGCCAGCCTCAGTTTCCTTGCCTGTAAAACAGAGGTGGGAATCCAGGCTATGCTTGCC
ACAAAAGGAGGACCAAATGACGACTCAGCATACTAGAAGGCTCTACAGCGCATATTATTACCTGTGTCTTTAATAATGTT
CTCGATTTGGTTCACTTTCTCCTCATACAAAGATGTGTATGTTTATGGAAAGAATATACAATGGATTCTGAGTGAACT
AAAATTGTCAACTGCACTGGGTATGAGTCTGCAAGCATGAAGGAGGCTGGGTGAGAAAGCCTCTTCAAACAGCATATGT
ATTCCTCTGGTGTCTGCTTTATTATTCACTTCTCTAATGTGAAAGTTTATGCTTTAGTGCAGAGCTCCTGGGAAATATG
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Fig. 6.32

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CAAAGAAAAAAGAAAAGATAACCTTGGAGGGGAAAAGAGGTAGTGCTTATGCCAATACAAAAGCATGAATAGAAGCAAA
TAGTTGCTGGTTTCTTTTAGCTGTAGTTCTTTAAGTGAATCTGGTATATATACAAGAAGGTAGTTAATGTTTTAAAGAT
GAAGGAAGGAATAGTTTCTTTTAATGAAAATAGACTCTAGTGCTACGGAAATGAGAAGTCCCCTGTGATGAGGCTAGAG
GAGAATTACTGAAGGATGAAGTGAACGTACGTGGCTCCATTTTGTTTTTCTTGGAGGCCATAAAGAAAACCAGACAGCT
TACTCAGTGTAAGTTATTTATACTTGTAAAAAAAATCACCCCATTTTTATAATTTAAGGGGATGCTAACGGTTCAGCTT
TGGATTGCAATTACATTAAATGTATTTTCTCATAAACTTGTTTTCAAATGTTTTAAAAAGATTTAACCAGAAAATCCAG
AATTCTTTTGCTGAATTTGTTTTTTAAATGCTTTAGATAAAATATTTTAGAATTATGACCTTGAATTTGAATTTAAGGG
ACTCCTTCAACCTCCTCTTTTAATCTCCTTTGTTATTTGGAGGGAATTATCCAAGTTGTNTCAGTTAGCTATTGTTGTG
TAACAAACAGTTCCAAAACATAGTTTTGGATTAAATGGATAAATGATTAAAAGGATAATCATGGAAGATTTTCATGAGT
CTACCAGTTGGCTGGGGAGTTCTGGTTATCTAAGCAGATTCTGATTATTTTAGCAAGCCTTACTCATGCATTAGGAGTT
AGCTGGCAAGCCAAGATGACCTTCCTTGGTAAAGTCAATGCTCCTTCCCATGTCTCTCATATCCCTCCAACAGGCTAGC
TCAGGGATGTTTTCACAGTGGTGTCTGGGGTCGAAGCATGGGCCAAGTCCAGTTGTGTAAAATGGTAGTTGGCAACACAC
ATGCACTTTTCAAGTCTCTTTGTGTGTCACATTTGCTAACACCCCACTGGCCAAAGCAGTTTGTGGTGGAACTCCGAGT
CCGAGTGGAGGATTGTAAAGTTATGAGTCAAAGTGTGCATACATAGGAAGGCCATGAGTTGGGCCATTAATACACTCAG
TCTACCAGAAGTCATTTCACCTTTTTTCCTCCATCTTAACAGCCTCCCTTTTGGAAATTTAACTTTTACATACTGTGTA
AAATCAAACCAAACAAACAGAAAAATCTCAACTTACCTGGTTACTTTTCAGCACTGTTTTTAAAGCTTGGTCAGGGAGT
AAGTTGTTATTACAGGCTAAAATTAAAATTTTGCTTATTTACTTACACACTTCTCCCCTAACTCCTCCTTTTTAAAGCA
AATTGAGTTTAAGCCACAAACTGTTGAATAATGAAGTTGGTTCTTTTTCAGTTTTATTCATATCTATCCCTGAGCTCTA
TTCTGCCTCTAATCAATTTCTGTATTTTGGAGCCGCTATTACTATAGCATTTGCTGGGTGAGTCCGTGGGGTTTTGATC
ATGCTTTCATATGTFCTTGCTGTTTGATTTCCTCTCTCCTTAACTCCAGGTTTGCATTTCTTCCTGCCTATTCTACATG
TCATAGAGGAACTTCAAACTGGACGTATTTTAAAGAATACTGGTTATCCACTTTGCCCAGTCACACATACCTTCAACCA
ACAAATATATATTTGTTGAGTGACTTCCATACTATTTCAGGCATTAGTAATATACTAGTGTAAAAGTAGACAAAATGGA
GGCCATCCTAGAGTTTATAATCTAATGGGGATTGACAGACATTAAACAAATGTGTGAATACATAATTTAATGCTTGGCC
GTGATAAGGGTTATTGAGAAAATAAAGCAAGGTGAGCAGAAAGGGAATGATGGCGGAGGTGCTTTTTTTTGATTAGTGTT
GAGGGAAGGCTTCTGAAGTGAGGGAGTGAGCCATGACAATCTGGGCAAAGTGTGCTGCACGCAGGGGCATTAGTGGATG
CAAGGCTGTCTGAATGACGTTTGGTGCTGCAGCAGCTGTACTTCCAATGGAAAGAAGGGTAGGAAATGAGGTGGGGGAG
ACAACAAGGGACCAGATCACGCAAGGCCTTTGTAGACAACAGTAAGCTCGTTACATTCTTTCTAAGTGCCATGGCAATC
CATTGGTGGATTTGGGGCAGGGGAGAGACATGGTCTGATTCTGGCTGCTGCGTGGGGCACAGGCTATAGGGAGCAAGGG
TGGAAGTTGGGAGGCTATGACAGTCATGGCAGGGGCTGTGTTAGGAGAGTAACAGTGGAGATTGTGAGAGGATGGCAGA
CTGTGGAGCTCTCCAAAATGTGTTTCATTTAAAGAAGCCAAATTTAGATACTGTGATCAGTTCTGCCAAGTCGAGCCTG
TGGAGAAGTATCTCCCAAAGTAGACCCAACAGACTTCCGGTGACATCTCAGTTACTTTTAGGTGCTATGTCACATGCAT
GAACTTTTAAATTTTAATAGTTACTGATGATTTTAATGTGGATTAGAAATATATAAAACTTGCATGTGAAAACTATGAT
TTTACAGTTATTACTTGGAATGAGGAAAAAGTATTGAGTGTAGTGCAGACTTGAATGCAGAAATAGCAAAAATCTCTAG
GGTTATATCTGAATGATTGGATTTAGAACATGCTGACAGGATTATTTGTTAGATTGAGTCCTGAGGGATAGTGAAGGAT
AACTCTGGTAAGTATGTAATCAGATTTAGTTATTACCCCTTATTGATTATCTACTGAATGTCACTCTATGTTAGGCACC
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TCATTTAACAGATGAGAGAAGTGAAGTTCATAGAGAATAAATAGCTTGTTTTAGGAAGTCTCTGATTTTTCAGATATTA
CAAGATTAGCTCTGGTAAATTCTCTGCAAACAGGAACTTAGTCTGTTTTGTACACTACTGTCTCCTTAGAACATAGAAA
CATTTCTAGTATATAATAGACATTAATCAATACTTGCTAAATGAATGAAAGGAAAAGAATGGTTCTTGTATTTAGATC
TTACTCAAGAGTGGTATATTTTGTTTTCCTTATTATAGAATGAGGTTTAGGAAGTAATCTCTTGGAGTTTAATGGAAAG
CCATTTTCGTCATAGGATGACTTCCTGTGTCCCCTAAGTGCAGGTTTCCCAATGTCCAGACCTTGGCCTTATATTTTTC
TCCTTATTCTCCTTGAAGGACCTGGCTCTTCTGGCTATAACTACATGTGCAATCCCCAAATCTCAGTGTTTGATCCTAA
GCTCCAGTTGTAGGATGTATAAACCCACTAACAAGCCTAGAGCCACTTTAGGGCAACATAACCTTGGCCAAGTGACTTA
ATTGCCCTTGAGCCTTAGTCTCTGTCAGTAGAAGGAAGAGATGGAAGAAGAAACTGAGCCTTGATTTGAACAAAACCTA
TGGGCACTAGAAGAAAGAAAGACAGGGTGCTTAGGGAGCATAACATTAGGGGGGAGAAGACCTAGAAAGCAAACAATTG
AATTTCTATTCCTTTTGAGCACCCTTATTTTGTTTTCTATCTAGAACTTTCTAAGGAGTTAAAAAAAGACTCTTTTACA
TAAGTTTTTTCAAAAGGCTTAAATTTATTTTTTCTTAAAAAAAGGTAAAGTATTGAAAATAAAGCCATTCAAGCATTTA
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TTGCTGGAAGGTCTGAGGCAGGGTGTGACAAACTGACCTGACTGAATGAAGGGGCTTCTGGAAAGGGGAGGAGCTCCAT
ACAGGAACATTGTTGGGGGTAAGGTGAGCCTGTGGAGGATGTCTCTCCTCTCTGCCTTTCCTTGCAGAGCTGGGAGCTG
TGGAGAGTGGAAAGTGTGTCTGAATGACTTCGCATGTTTAGTGAAGCTCATTGCCAGAACTGCGACTTCCCCTGTTGCC
TTGAATCTGCTGATCAGCCCTGGCAGCACACGTTTTTTAAATTATTTAAAAAGGGACAAGGGTTATAAAGATGAAAACC
ACATTTTTACCAGCCTCTGCAAACTCCTAATTTGGTGGGTGAAGGACACATCAATGAAATTTCCAGTATAAGGAGTGGA
GGTAAGTGCTATAGTCTGAATTTTTGTATCCCTCTCAAATTCATATGCTGAAACCTAATCACCAAAGTGGTGGTATTAA
GAGGGGGTTTGGGAGGTGATTAGGTTGTGAGGGCTCTCTCTCCATGAATGAATTAGTGCCCTTATAAAGAGTCTTGAGG
GAGCCTGTTTGCCCCTTCCATCATGTGAGGACACATAGAAGGTGCTGTCTATAAGGTATGAGCCCTCACCAGACATTTA
ATCTGCTGGTGCCTTGATTTTGAACTTCACAGCTCTCAGAACTATAAGCAATAAATTTCTGTTGTTTATAAATTGCCCA
GTCTAAGGTTTTTGTTATATATATATATATATATATATATATATATATATATATATACACATATATATATATGGCAG
CTCAAAGGGACTGAGACAGTAAGCCAGACACAGAGGTCAACACCTGCCTTGGGAAGATAAAAAAATTCCAACTGGGTGG
ATCTGAGTGATGACTTGGAAAGAATAGGATTCAAATTAGGCCTTGAGGCAAGGATGTGACTGATGCTTATCTTTAGGAA
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Fig. 6.33

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AAAAAGGTTGTTTTACTCTAGGTAAAGGAAGTAGCCTCTTCAAAGGCTTTGAAGCAGAATAATGCGGCGGGCGGGGGCT
GGGGTGGGGTCAGGGGGCATGGTACATCAAACAGTGGGGAGGGTGTAGAGTGCATAGTGTTGGGAAGAACACCATGAAA
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GGGAGCCATTGACACTTTGGTGAGAGGATCTGAGGAGTGGAGTGGCAGGGGAAGGCCATCATCAGCTCTGTGTGTGCCA
GGGCAGCAAGATTTGTTAGCTTAAAAAAACTTTTTGAAATAATTTTAATTTAGAGGAAGTTGCAAGAATTGTCAAAAAG
TTTTAGGATATTCTTCATCTGAATTACCAATTGTTAACATTTTGCCACATTTACTTTATCATTCTCTTGTTCTTTAAC
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TAATATCTAATATCTGAAGTCTGATCTCCAAACTTTATTCAAATTTCACCAGTTGTCCCAAAATATAGGAGGTCTCTTT
CTCCCTTTCTTTCTTCCTTCCTCCCTTGCTTCTCTGCCTCCCTTCCTTCCCTTCCTTCCTTCCTTTCCTTCCTTCCTTC
TTTCCTTCTTTCTTTCTCTTTNTTTTCTTTCTTTCTTTTTTCTTTCTTTTTCCTTCCTTCCTTCCTTCCTTCCTTCCTT
CCTTCCTTCCTTCCTTCCTTCCTTCCTTTCTTTCTCTCTCTCTCTTTCTTTCTTTCTTTCATCTCTCTCTCTGTCTC
TCCCAATAAACTAATCATAAGTTGAAAATGCATTTAATATGCCTAACCTATCAAATGTCATAGCTTAGCCTAGCCTACC
TCGAATGTGTTCAGAGCACTTACATTAGCTTGCAGTTGGGGAAAATCACCTAATGTAAATCATGTAATAAAGTAAATTA
CATAAATATTTCATGTAATTTATTGAATACTGTACTGAAAGTGAAAAAAGTATCCAAAATATGGTTTCTACCAAATGTG
TTTACTTTTGCAGCATCACAAAGTTGAAAAATCCTAAGTTGAACCATCCTAAGTCAGAGACCATTTGTAGCCTAGGTGA
CAATGAAATAGGCTTGAGTGGAAGCACTGACAGTGAGAATAAGACAGCCATTTCAGAGTTGAACACAATAAGATTTGGC
AACAATGGGCTTTAGGGTAAGAGAGGGAGTAAAGCTGGTAATGAGGTTGTAGTTTACAGGATTGATGTCCTTAACCAAC
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TCTATAGTGCTTTCAAAGGCAGAAAGGCAGTATGATTGGAGGCAGCTTATGGCAGTGATCAAGAGTAGGAACTCCGGTG
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AATTGGGACTCAGGTGTTACATGGTTTGCCCAAAACCATATAACTATCGGTAAAAATGCTCATTCTGTTCCACCTACCA
GACTATCCCCGGGCTCGCTACATGGGGATACCTAATAGGCCATAGTAATGGGCTAGTGTCTCAGCATAGGCATACTTTG
GGAGTATTGTGGGTTCAGTTTCAGACCACTGCAATAAAGTGAATATTGCAATAGGGCAAGCCATGCAAACATTTTGGTT
TCCCAATGTATGTAAGTTACATTTCTAGTATACTGTAGTTGATTAAGTGTGCAAATAGCATTATATCTAAAAAAAGTAC
ATACCTTAATTTAAAAATACTGGTTGTTAAAAAAATGCTAATGATCACTGAGCCTGCAGCAAGTTGTAATTTTTTTGCTG
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CATCTCAGGAAACCACTTTCTTGGCTTATGTGTAAGAAGCAGCTCCTCAAATCTTCAAATTTTATCATGGGATTGCAGA
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GCCTCCTCCCATGAGTCATGAATGTTTTTAATGGCATCTAGAATGGTGAATTCTTTCCAGAAGGCTTTCAACTGACATT
GCTCATATCCATTAGAGGAATCACTGTTTATGGCAGCTACAGCCTTACAAAAATATATTTCTTAAATAATAAGCCTTGA
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GTTGTTTAGTGTAGCTACCTTCATTGAAAATCTGTTGTTCAGTGTAGCTACCTTCATTGTTATCTTAGCTATATTATC
CAGTTCACCTGCTGCAACTTCTTCCTCAGCATTTGCTGCTTCACCTTGTACTTTTCTGTTATGGAAACAGTTTATTTTA
TTAAACTTCATGAATCAATCTCTGTTAGCTTCAAACTTTTCTTCTACATCTTCCTTACCTCTCTCAGCCCTTGTAGAAT
TGAAGAGTTAGGGCCTTGCTCTGGATTAAGCTGTGGTTTAAGGAAATGTTGTCACTGGTTTGATCTTCTATTCAGACCG
CCCAAACTTTCCCTATATGAGCAATAAGGCTGTTTTGCTTTCTTATCATTCATGTATTCACTGGAGTAGCACTTTTAAT
TTCTTTCAGAACTTTTCTCTTACATTCACACCTTGGCTAACTTGCACAGGAGCTTTTGGCTGTTCTTGGCTTTTAACCT
GCCCTCCTCACTAAGCTTATTCATTCCGAGCTTTTGATTTAAAGTGAGATATGTAACTGTTTATTTCACTTCAACACTT
CAAGGAAATTGTAGGGTTATTAATTGGCCTAACTTTTAATATTGTTGTGTCTCAGGGAGAAGGAGACCTGAGGAGAGGA
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GGCATGGTTTGTGGCATACCAAAACAATTACAACAGTAGCATCTGAGATCACTGATCACATATCACCCTAACAAATATA
ATAACAATGAAAAATTTTAAATATTATAAGAATTACCAAAATGTGACAGAGACATGAAATGAGCACACATTGTTGGAAA
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TGAAGATGAAGGGGAGCTGGTGACTGAGACTGGGTAGGAGCAGTCAGAGAGGTAGGAGACAGAACAGGAGGGGGTGGAG
TGAAGGAAAGGAAGAAAAGTAAGGGCTTAAGAAGGACTTGGCCATTCTTTACACAAACAAGGCAGACAAATTCAGAGCA
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Fig. 6.34

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GAGACCTGGGATAGTGATGAGTGGGTTTCTTAGTGACCATAGTGAGAATGGTTCATTGGCTTGGTGAGGACAGATGCTT
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AAAAGGAAGGAAAACTTTGAAAGAAACAGGGTAAAGAAATGATCTGTCTCCAAAAGAGGCAATTGTCTTGGTCATCCAG
GAGGTAAAGGCAAAGGCTTGAAGATGCAGGAAATGGTGGGGACGATGGAGAGAGCTGGCAGGGATAGAATCAGGAGCAT
GGCTGGAGGGATTGGTCTCAGGAAGAAATAGAGACACTGCTTCAAGATTGGAGGGAAGGCTGGGCACAGTGGGTGGCTC
ATGCCTGTAGTCCCAGCAATTTTGGAGGCTGAAGTAGGAGGATTGCTTGAGTCCAGGAGTTCAAGACCAGCCTGGGCAA
CATAGTAAGATCCCCATCTTTGCAAAAAGGAAAAAAAAGTTAGCCAGACATGGTGGCACATGTTTATACTTCCAGCTAC
TTAAAAAGTAACCCTGACTCTTAAAAAAAAAAACTGAAGGCAAGAAAGAGACTGTGTGTGATGACCTATGACATTTCAT
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GAAAGTTAGAAGTATATTTGAGAATCTGAAAGGGCTTGAAAAGTTAGGATGTAGACTGGGCATGGTGGCTCATGCCTGT
AATCCCTGCATTTTGGGATGCTGAGGCAGGTGGGTTGTTTGAACTCAGGAGTTCAAGACCATCCTGGGCAACATGGTAA
AATCCCATCTTTATTAAAAATATAGAAATTAGCCAGGTGTGGTGGCACTCACCTGTGGTTCCAGGTACTCAGGAGGCTG
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TTCTCACGAGATCTGAGTATTTAAAAATGTGTAGCACATCCCCTTTCTCTCTCCTGCTCTGCCATGCCAAGACGTGC
TTGCTTCCCCTTCACCTTCCATCACAATCATAAGTTTCCTGAGACCCCCAAGTCATGCTTCCTGTTAAGCCTGGGGAGC
TGGGAGTTAATCGGACCTCTTTTCTTCATAAATTACCCAGTCTCAGGTAGTTCTTTATAGCAGTGTGAAAACGGACTAA
TAGAGAAAATGTGACACCCCAGAATTAAGGCAGAAGAAATGGATTGGGCCAGTGTAAGNATGGCTTTTAGTGAAAGGTT
AAGAATACTAGGAAATCTCAGAGTTCAGAGAGAATTCAGTTGGAGATGGTTGACCATAGGCTGTGAAGGCAAGAGGAGG
CTGAACAAGGTTCCTGTGAAATATCCATAATGACAGAGAAGAAGGAAAGCACATTTCTGGGGCTCCAAGTAGCCTCTTA
GAAGTTGGAAAATTCTCAAGTCTTAAGTTTTAACTTTAAAATATTTAAAATTCCAGGGCTGGGCCTGGTGGCTCATGCC
TGTAATCCCAGCACTTTGGGAGGCTAAGGCAGGCAGATCACAAGGTAAAGAGTTCAAGACCAGCCTGGCGAACATGGTG
AAACCCCGTCTCTACTAAGAATACAAAAATTAGCTGGGTGTGGTGGTGTGTGCCTGTAATCACAACTACTCGGGAGGCT
GAGGCAGGAGAATAGCTTGAACCCAGGAGGTGGAGGTTGCAGTGAGCAGAGATCACGCCACTGCACTCCAGCCTGGGCA
ACAAAGCAAAAATCCATCTCAAAAATAAATAAAATAAAATAAAATAAAATAAAAAATATTTAAAATTCCAGTGCNGAT
CTGTTAAAATAAAGTAGTAGATGCTAGTAGGTGATTGATTTCTCAGTAAATAGACACCTAAAATATATATGCCATATTT
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GAAAAATTTGAGACTGGCAGATCTTGAAAATGAGACAATCCTAGTTGATAATGAGCTCCAAAGTGTAATCAAGAAATCT
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TACAAAATTTCTCTCTTGTGGTGCACAGAAAAAATATTAATGTTTAAAAAGATGCATTTGTAAAATGTTAAGCCAGACA
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TGAACAACAAGCAGCCAATTTGTGGAATTAGTGGCTGTTTCAGGAGGGTAAGTAACCTCTCATTTATGACCATCTCGTT
CCTATGTTTTAAAATAAGATTGTGGACAAATACTTCCATTCAACCTAGTAGATGCATTTTGGCAATGGCAGATCACCCA
AACAATGGATAACAATAGCAAAAGAAATGATATAGTCAGAAGTCTAATGTTAGCCATTTGTTCCTAGAGTTTCGAAATA
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TTTTTTTTTTCACTTTCACTGCAGCATTTAAAATTACAAGAGATATTCAGTACTTTATTATAAAGTAGGTTTTGTGTTA
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GATAGGTATATTAAATGCATTTTTGAATTAATGATATTTCAACTTACAATGGGTTTATTAGGACATAATCCCACTGTAA
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CTTGGACCATCTTAAATGAACCAATGGGAATGTCACCATGCACATCCTTCCAAACTTTGTTTCAAGCTATTTCATGTCA
AATTTATGCTTATACAAGTGGAATAATATGTATGTTATGTCTTATGTTTTAAGGAAGTCTTGACTTATGGAGGGTGAGC
ATTAGTCTCACTTTTAAACTAACTTAGGAAAATGATCTAAAGCATAGTGAAAAGAACAATAGAGCAAGAGGAATGAGG
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TTCAGGAAGCAAGCGGGTGCATTTAATAAATTAAATTTTTATTTAATTATTTTATTTTTTATTTATTTATTTTTTAAAGA
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AATCCTCCCGCCTCAGCCTCCCAAATAGCTGGGACTACAGGCATGTGCCACCATGCCTAGCTAATGAAAAAACAAAAAA
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CCTCCGAAAGTACTNGGATTACAGGTGTGAGCTGCCATGCCTGGCCTGGCCTTGCTTTTTTATTTTTTAGCTTTTATAG
CTCCATATCCTCTAATGCCACCACAGTATGATGAGTGGAAATTTGGGTACCATGTTTAGCACTGATTTTCTCTCTTGGC
GAGGAGGCACCACTATCACGCTATGTTGAGTTTTTGGCAGCCAAGATCAGAGGGACTGTCTTGTAATGGCCCCATCCCT
GAGATCTGTGGGAATAACACAGGAGACAGGCACTGACTGCTTCCAGGGGAAGCAACAATCAGTGCACTGAATGCCTTAC
CAAGCTGAGTATCGCTGAACTTCAAAACAGTGCCAGAGGAGTAGAGTTTTTTTAACTTCCTCTGCTCTGTCAACAGCTG
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Fig. 6. 35

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ACTTAGCTGATTTTTGCCTGTGATGATAATATCCAGAGTGATGACTGGATTCTCATAATCTGCTTCTCTACTGTTAAGG
AGGGATTTTTACCTGATAGCATTTCTCTAATTTTACTTATTTTATTAATTCTTCTATGCCAATTACTTATAAATTTTGG
CTTATTTCTGATCACACTATACCATTTGTAAAACATGAAATTCAAGTTGATTGATTGCTATAATTTTCTCCCATCCATA
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TTACATGGCTTCCTGGAGACATTGATTTGTTCTTCTGGAAGGGGTTGGACCTCCATTGAGAATCACTAATTTAACAAAAA
GCCCACAGATCATTGTATGAGTGTGTGTGTGGATGGGGGAGGGGGAGAGAGAGAGAAAGAAGGAAAGAAATGGGTTAAAATTT
CATTTTAAGGAATCATTTTATTGTTGGATTAGAAACTAAAGACTGTCCATAAAAAAGAGGAACCATTCTAAAATTTTCA
TAACTTGCTGTATGTGTGTATTTTATTTCTTAATAAAGCTCAAGTAGTTTCTCATGAAAGTCAATGGATAGAGAGCCAT
TTCAGGACNTGTTTGAATTGGCAGTTGGTTGGAGTTTGGCCATGGCAGGAAAGGACAGCTGTTTGTTGTGGGGCCTTGT
CTCCATGGTGTTGGGCAGCACATCTTCAGCTCTTTCTGGGAAGACCTCAAAGTAGGAATAGACCAAGAAAAACAGCAAG
AGGGCATGACTGCATGCTGGTCAGATGGAATTCAGCCTGTACTGCTCTAGCAGAGCTTCTCCAGGGGTTAGAATGTTTT
GCTTTTATAGTCATCTGGATTGTTTGTTTACTAAGGGTAGAAAGGTCCCAGCCTCACCCAAGACCTCCCGATTCAGTCT
CTGTGGGGAGCTGCTTGGGAATCTGTATGTTAGAGAAGCTCCCAGGATAATTGTGATGTGTGGTCATGTTTAAGAGCTG
CTAGTGTAAATAATGGTGAACAGTAGCACTGGCTGGAAGCACTGTAGAAATTGTGTGTGTGTGTCTGTGTGTGTC
TGTGTGTATTGTGCAAGAGCTATGAGATAAGATCAGGCTACTGGAATTTGCCAAGAAGCAGCACGGGAGACACATTAAC
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AAAAGATTGAGCAAACTTTGTAGGAAAACTACAGAAGTTGGAACCACATATGATGCTTTCTATTATATGCCCAAAGTCT
GTTGTATATGATGAAAATGAGGTTTTATTTACTTCTTGGTACACCAAACACAAACACTCTCTAATGAGGAAAAGAACCC
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TTAAGGGAAGACACTATTCTACAGTTCGGTGTAGCTGGGATGACTTTATATCCTAGTTTGCAAGACAGTCCTGGCCTCA
CCTGTTGAGCTGGTGTCCCATCTAGTTAAGATACCTTTCATGCTCCCCAGTGTCCTGGTTGAGATACTCCGTTTATGTG
GTCACCGTGAGGATAGGAAACAGAGAGAGGAGATTTTTCTTTGGATCACTGAATAGTACTCATTTCATTATATGATTTA
TCCCATTCCAGGCTCACCCCTAACATTTGAGGGCCCAGGACAAGACCCCCCTATTCCACAACCTATTCCTTCTCTTCTT
GCCCTGGCTCCACTGTGCAACTGGAGGATCCTTGTAAGAATGTGTGTGGGTGAGCCAGCCTGCTCATTCAACTTTATTC
ACACCCACATCCACTAGCTGCCCATGAACACCCCTCAGGCCTAGGGGTGCGCACACCTGTGAGTTGATCTGCTTCGGGA
GGGTGAAGACACACCCAGGCCCTGGAAGCTGGTTTCGGACCACTTGGGGCAGGAAATTCAAGGGTCATGAGTACTTGGG
GAAGAGTCTAGAAGGGAAGTTGTNCCTGCAGGTGGGTATGTATCACAGCCCTGCAGACTGCTGGTTCCATGGAGAGGCA
TGGCTGGAGAAGGGCAGGAGAGGGCATTCTGAAGCATCAGCCCCCTGATTAGGGATACCTGGCCCTAGAGTGGTCCTGG
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TCTGAAGTGTTTGGGTTAAGAACGTTTCTCATGTGAATAATGCTTGGTCACCAGACTGAGGTCCAGCGTATTCTTCATC
TGGTCTGATAGAAATCCCTTGTCTTTAAATGTCATATATTTAAGGTAACAACTCTATTTTGTAGGTTGAAACAAAAACT
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GGTGAATGTTTTTTTGTTTGCTGGTATGTTCCACCATGTGTGTGAGCATGAAGGTAGGAAAACTGTTTGCTCAGCCATT
CCCTCTGTGGTTTGGTTTGTTGGAGAATCAGTCAAATGGAAAGGCTTTCCTCTGATGTGATTACAGAATCAAATATTTT
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GCCTGATTTTCCTCCTCTTTCTGTCACCCATCATAAGCATATGGAGAAGATGGATGTTTGAAGGGAAAATAATGTTATT
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TGTTAAAGAACTAACTATTATATAATAAATTATCAAACAATTACGCCAAAGGTATGTTTACAAACTCCTGTTCTATCTT
GGTGGGTTGATTAGCTTCCTTGGGGGGTCCAAACAGCTATGGCTTTTCCATTTTTATAAATGGTTCTGCATTTATAAAA
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TGAAATGTTCATCTGGAAAGGATTTGATGAAATGTTCAAGTCAAATGCAGAATAAAAGCCAGCAGGAAGAGGCAGCATGA
GGAACCAGAGAGCTTCTTTACCTGGTTTACTGACCATTCTGATGCAGGTGCTGATGAGTTATGAGAGGTCACCAAAGAT
GATATTTGGTCAAACTCATTACAGTACTACTTGGTTCCCGATATGGATGTGAAGAAGCAGAATGAGAAGAAGATGATG
ATGATGAAGAGGAGGAAGGATAAAAAGATACTAATGAAGAAGACAATGAAGATGAAGGTGAAGATGAAGATGATGA
TGAAGGGGAGGAAGGAGAGAAGGATGAAGGAAAAGATTACTAGAACACTGATAGGTTCCAACTTTCCTGTTTAAAAATT
TTTCTCCAGTCCCTGGGAGCAAGTTCCAGTCTTTTCTTTTTCCTCTTGTGCTCTGTTGCCCTGTTGTTGAGGTCTCTTT
```

Fig. 6.36

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GCTCTACACCATGGTTCTCAACTTATTGCAGAATACAATGGGAAAAGTGTCTCTACGCCCTTTCTGTTTGAAATTCATTT
TTATCCCTTTCTGTCTGAACAAAAACTGTATGGAATCAACACCACCGAGCTCTGTGGGAAAAAAGAAAAACCTGCTCCT
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GTATCTTGGGCTTAGAGAGTACACGGTGTCTCTATGTGAATATGGACAGTTAGCATTTACCAACATGTATCTGTCTATT
TTCTCTTGTTTAAAAAAAGAAAAAAACTAAAAAACAAAATGGGATTATAGAAGGTCAGCAAAGGGTGGATCTGAGATGT
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AACCTGTAGGATCTTATTTGGAATTGACTTTCTCTATTGTAATTTTGTTCCTGCTTATTTTTAAGTTTTCTTTTTTGTTT
CACTGTAAAGGAAAGATGATGCTCAGTTTTAAACGTGAAAAGTACAAGTTGCTTTGTTACAATAAAACTAAATGTATAC
ACATACACACACACACAACACACACACACACACACAAACTTATACCAAAATCACACCACTTTAGTTAGTTGTCCA
CTGGAAAGGTGCTCTTTGTTTTCGAGCTTCAAAGAAGCTCGTCTTCTTTTATTTGACTGTCGCTAGGGACTCCACTGAG
CAGGAATGACTATTCTGATATTTGCTTGTTACATTTTTGGTTTTCAGAGGTGGCATATTGAGGAGGTGCACTTATGGAT
TGCAGCATAAGTATCCATGAGTCCGGGTGTGGTGCGTGTGCTCACTCATTATCTTGATGTGCTGCTCACTACCACGTCT
CTCCTTTACACTATTGGAAACTTTCATTCTTAATAATTTAATTTATCCCTAGTGACTCCAGCCTCCCATGACAGACTAA
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TGACTTATGTCTTGAAAAAACTTTAACTTTGGGAATTTTGATGTATAATGAAGCCAGGAGAGGGAGACGTGTCATTCCA
AAATGCTGCGAATAGAATTTAAGCAAGCAACATGACACACATTCTTTGTTTCTTTTGCTTTTTTTTTTTTTTTTTTTTTTT
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CTCTCTAAGGTATATTATTGCCTACTATATCCTGTGCCCTGTGCTGGAATTTCTCTTGTGCTAATTTTTATTGATAACA
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Fig. 6(37)

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ACCCTTTCCCTGCCTTGTCTCTGGGTTTTTGTCTCTCTTGCCTCATTTTTCCTTCTTTCTCTCTCACTTTCTTCCCCCA
CCCCTTCTCTCTTGTTTACACTCCCCGCTCATGTATGTGCCGCTTCGTCTTTGTGTTTATATCTACCAGGGGAAAGACC
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Fig. 6.38

ATATATATGAGCTAAATATGGGGTTGAATCCCCACCAATCATATCTTAACAGATTTTTTTAAAAAGGTAAGTATCTGAA
TCAGCTGCTTTATTTTATTTGTCTCTTTCTCATAAAAATGACAATAAAATTATGGTATGTGAAC CTGATTATTTAGA
TTCCAGTTAAACAGGATTTTACTTTAGTTTCTTTTTTTTTTTTTACTGAAAAAAAATTTTATTGACTTGTTTTTTGGTG
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CCTCAAACTCCCACATATGACTTCTTTTTTTTTTCTTTTTATTTATTTATTATTATCATACTTTAAGTTTTAGGGTACAT
GTGCACAATGTGCAGGTTAGTTACATATGTATACATACCATTTGACGCAGCCATCCCGTCACTGGGTATATACCCAAAG
GACTACAAATCATACTGCCATAAAGCCAGGAGAATTTCTTGAGCCCAGGAGTTTAAGACCAAGCTTGGCAATATAGCAA
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ATCAAATGTCAGAGATTTGTATTCTTGGTAGAAACTGAGATTTCATTTTCCTTCATCCCATTCTGATGCTGGCTTCCCT
GAAGCAACACTCCTGTTCCAGATTTGTATGCTTGAGGTAAGTCAGAAAAGGAATTTCGGTGAGGCATGAGTTTGTTGAT
GAAACGTGGGGGAAAACCAAATGGACATCTTCAGAATGAAAGTGTACGTCTTTGTCTGTCTGAATGTGTGAGAAAGAAG
AACACTGAACACAGTCTAAACCTCCTGCTGGTGTGATTATTGAGTAGTGTTGATGGCCAAAATATGAATGAAGGTTTTT
CTGTTGCAAAATGCTTTATGTTATGTGCAGGGCAGACATCTGTGTGCCTGCCTTAAATTTATGAACTAATAACATTCTG
TTCCAAGAAATCCTGTCAGGTCTTTAACGGTACCATTTTGTTCTGAGTTATTCTGGTAAATGAATGATTTTGCTTTTAA
GTAATTATTTTTTGAATTTTTAAATATAAAGTCACATATGGTTTAATGTATGGAGAGGATGCCTTTATCATCAATGAAA
ACTTTAAGAATTGACTGATTGCATTTTATTTTAACAAAATATCATTCATTATTTTCTTCTATTTGGAACAGATGTAAAT
GTTCCAATCTATTTTGGATTCAAATGAATTTTGAATCAAATAGATTTAAAATGAATCTATTTTGAATCAACAAAATAGA
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TCTTCTGATGAATCTCTTGTCTGTAGTTTTGAAAGGAGGGGAGAAGCAAAGGAATTAGCAATCACAAAGATAAACCTGC
AAAGGCAAACTAAGTAGAAACATGTTTTTCCGTTGCAAATTATGCTTAACTACTATAGGCTGTAATCCCTATGCTGACA
TATCTCAGAATAAAATGTAAGTTGCTTGAGATGCTATTCTGAAGCATGTGATATGGTTTTATTGATTGATAAATTTATC
CATATGATGATAGTTGTAGATTTTCAGAGGCCCCATATGGACCATTTGAAATGAAATGTTATATTGGCAATGGATGAG
AACATGCCACTTGCCCTGCTCCATACAAACTTTGATTCCTAGCAATCAGCACATGTTAGAGAGCTGGAGGCCACGGTAT
GGTGTGCAAGTGCTCCTGTGCTGAAGTGACACCAGAACAATCACCACATTGCCATCAGGCCGTCCATCTGAGGTAGGGT
TTTCCTTCTTGTTTCAATCTTTATCATTCCTGGAGAATTTGCCAAGGCCCCCAGGGCAGTGCTGAGGCCTCCAGCGCCC
CCTACAAATCTCTTTTTCTGTCCCTTCAACATTTTCAGGTCTAACTACCTGTAGTTTTCCTCTCTAATTTCTGGGCCAA
AGCTTGGCAGGGGGCAGGCGCAGAATTAGGTTGAAGAGGCCAGTATGAGCAGGCATACCAAGGTCCATTGTGCTATACC
AGAGGTGTTACATTAGACTAATGAAGGGATTATAGAAACTTAGCTCTCCATCTGAATTCTCTTATTAACACATCACTGT
GAATCAGCCTGCTGCTTTCGGTGTTGTGTATTTTTTCCCTTTGTTACTTCTGTTACTGCTAACGATATTGGTATCATTT
GTTGTATCAATGCAACTGACTTGTACTTCAAATTAATATCTCAACCAACCTAGAATTTCTTGTCTAGCTATGGTAGGTA
GTATGGGGTGGTAGTGGTGTTGTCTCAGGCTAGTATATGATAATATGATTTGAAGTCTGTTTCCCCCCCATTTAACG
AACTAGTGTGAACATCTATCCATGCCTTTTAAACACTTTTCTCCAAGATAATTTTTTATAGTTTTTTAATGGTTAAATA
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TATTATTAACTATGGGCATAATGTTGCCCAGCAAATCTCTAGGACTTAATCATTTTACATAACTGGAACTTTATACCCT
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Fig. 6.39

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TTATTTTACATGCAAATCTTAATCCATCTGGAATT..TTTTGTTGTATAGGAGTGCAAGAATTTAACTTTATATTTTTC
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AGCAGGTCCTCCCCTGGTTGACATTTCTTTCTTACCTGCCTGCCCTCTCCCTGACCACTCTTCCACATCCCCACCCTTC
CCCTCCTGGACTGCTTGCAATTGTCTATCCAGTCTGCTGTTTCACACCCCAGTGCCTTTTTACATGTTATTCCTTCTGC
CAAAGTGTTCTCCCACACATACCTCGCCCATCTGGAAATGGTTCCTTGTCCTTCAAACAACAATCAAAGAGCCTCCTC
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TGGCAGGACATAGCCCTGGTGAAATCACTGGCAGGGGCTTGTGTGAAGTAACTGCATAAGAAAAAAACAGCATGCTGTT
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CCTGGGGTGTTTATAACATGGAGGACCTATTTCCTGAGCTACCAGAGTCCTAACACATACCTAATTATGAGTTGGTCAC
CAAGAGTTATCAGTAATCACAAGTAATGAAAGCATGGCTTTGGGTGAGTGGGTGGGTAGGGTGGGTGAGGTGGGATATG
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GTCTTACTCTGGGCAAATGAACCTTGTTTTTTTGAGAATGTTAAGTTCATCTGACATAAGCCGCTCAATTTCACTCCTC
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GTGCTTTAGTTGCCTCGGTGAAAAATGGGGATAATCATAGTGCTGCCTTATAGGGTTGTTGTAATAATTAAATGATTAT
```

Fig. 6.40

CCTGCACATAGTAAACAGTCAATGAATTTATGCTATTATTATTAGTCTGCTATTTGTGGGCTTTCATATTTTGTCTCAA
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GCCTTTGAAGGGTTTTAAGCTGGGGAATAAAAAGATAATACTTTACTTTTAAAATTGCACCCAGGATGGATGGAAGACA
GAGAGTGGGAAGATGAGTGAAAGCAGAGAAAACAGTTGGGCACTCTTGTAGGTGAGAGACTGCATGGCCGGGGCTCAAC
TGGCAACAGTGGAGATGGAGAGAGGTGGGTGGAATTGAGATTTATTTTGAAAGTAGAATGGACAGGATTTGCTGAAGAA
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CATTCTTCAACCTCTACAATTATGAGAAAGTAATGACTGTCAAAAAAAAAAATTGCTCATTTTTTCTGACATCTTTCAAG
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CAAATAGCCTGCCCAAAACAACCAGTTAATAAATGGAATTGGGATTTAAACTTAGATTTGTCTGATTCCAAAGCTTGTG
CCCAACCAGTGTTTATTGAGTACCTTCCAAAGAGCCAAGCAACAGTTTGAAATAGGTTCACATATGTTCAACTATCACA
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AGATGAATTTGCCTACCTAAAGAAATGCTACCTGAAAAGCTTCTCTGGCAGGCTTCTCCAGAGTTTCATTATCTACTCT

Fig. 6.47

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GAACTTTTGTTTACATAGCAATCATTTCTGCTGTGTTCCTTTTCCCTACTAGCCTTGGTAGGCTTCTTAGCTGAATTGT
CCTGAAACCTATAAACCAGCCCTGGAGGCTCTGAAAAATTAAAGATCTGTCTTTTTTGTTCTGTTTAATAAATGTTAGC
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CAAGTTGCAAATAGAAAGAGGTATTCCCCAATTAGAAATAAAATATTGGGCTTCAGTTTCAAAAAAAAATCTTGAAAGC
TACATGCTCCATAGACTTCTAAGCAAAGAGAGTCAAGAGACAATCATCTTCATAACTAACACCAATTGTCATAAAGCTT
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GGAAGATTTAAACAATTTAAATATATGCTCTAGTTGTTTGATTTAAAATGCTTTTTGTGCCAAAGAAATTCAGGATAGA
GATTTAGTAATCAGAGTTGAAAAATGCATAACACATTGTTCTAGTAATTCCCATCATTCAAAAGGAACCATCTGTACTG
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CAAAGCAAAGACATAACAACAACGTAAAAACTTCAGACCAATATCCTTGATGAACATAGATGCAAAAATCCTTAACAAA
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GGGAGAGCCTCTGCCCCGCCGCCCCGTCTGGGATGTGAGGAGCGCCTCTACCCGGCCGCGAACCCGTCTGGGAGGTGAG
GAGCGTCTCTGCCCGGCCGCCCCATCTGAGAAGTGAGGAGACCCTCTGCCTGGCAACCGCCCCGTCTGAGAAGTGAGGA
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Fig. 6.(42)

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GCCCCTCCGCCCGGCAGCTGCCCCGTCTGAGAAGTGAGGAGCCCCTCCGCCTGGCAGCCACCCCGTCTGGGAAGTGAGG
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TAAGAGTCATCACCACTCCCTAATCTCAAGTACCCAGGGACGCAAACACTGTGGAAGGCCGCAGGGACCTCTGCCTAGG
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CTGCGAGAAACACCCAAGAATGATCAATAAAAAAAAAAAAAACAAACAAACAAAAAACAAAACACAAACAAAAAAGAAAG
GTTTGAGCTAAGAACTTGCAGGAGACAAGGAAATTAGTCAAGCAGAAGGATATCTGGGGGAATGGCATGCGAGGCAGAA
GGGAAAGCTAGGGTCGAGGCCCTCAGGGAAAGAAGCAAGGCCAAGGGGCTGGAGTAGAGGGAGGAAGAGGGGAAGTAGT
GGAAGATGAGACTAGCTTTACTACTGATTATGATGTAAGAATAGTGGCCAGTTTCCTTTCCAACTTGGGCCCGGCAGAA
TGGCTCCTGCAAAGAAGGGTGATGAGAAGAAGAAGGGTCATTCGCCATCAACGAGATGGTGACCCGAGAATATCCCATC
AACATTCATAAGTGCATTCATGGAGTGGGCTTCAAGAAGCGTGCCCCTCAGGCACTCAAAGAGCTCCGGAAACTTGCCC
TGAAGGAGATGGGAACTCCAGATGCACACTTTGATACCAGGCTCAACAAAGCTGTCTGGGCCAAAGGAATAAGCAACGT
CTCATACTGTATCCATGTTCGGTTGTCCAGAAAATGTAATGAAGATAAAGATTTACCCAAACAAGCTCTATACTTTGGTT
GCCTACGTACCTGTTACCACTTTAAAAAAATCTACAGTCGGTGTGAATGTGAACTAACTGCTAATCATCAAATATACCA
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GGAGTATTCCTCTTGAGAACTGGAACAAGACAAGGATGCCCACTCTCACTGCTCCTATTTAACATAGTATTGGAAGTCC
TAGCCAGAGCAATAAGGGAAGAGAAAGAAATAAGAGGCATCCAAATAGGAAGAGAGAAAATCAAACTACCTCTGTCAAA
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CATAACATTTATCATACAAATTTGGCATTTTTTGAGGGTGAATGAATGGGAGAACTATTAGTAGTCACATTGGGACAAT
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TATCAAGGACTTTGGACCGAAGTCTGTCTGGTTCCAAATTCTGGTTCTCAGCATGACTTTGAGTAGATTATTATATCCT·
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CTTTCTAGGATTCCAATTACATTTACGTTAAGACTATTTGGTATTGTCCTACCTATTCCTGAATCTTAGTTCATCTTCC
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AGGTTTGTTACATATGTATACATGTGCCATGTTGGTTTGCTTCACCCATCAACTCATCATTTACATTAGGTATCTCTCC
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Fig. 6.153

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TAATGCTATCCATCCCCCACCCCCCACACCCCCCAACAGGCCCTGGTGTGTGATGTTCGCCACCCTGTGTCCAAGTGT
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GGCTGGGTCAAATGGTATTTCTAGTTCCAGATCCTTGAGGAAACGCCACACTGTCTTCCACAATGGTTGAACTAATTTA
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GCCATTCTAACTGGTGTGAGATGATATCTCATTGTGGTTTTGATTTGCATTTCTCTGATGACCAGTGATGATGAGCATT
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GTTGTTTGTTTTTTTCTTGTAAACTTGTTTAAGTTCTTTGTAGATTCTGGATACTAGCACTTTATCAGATGGGTAGATT
GCAAAAATTTTCTCCCATTCTGTAGGTTGCCTGTTCACTCGATGGTAGTTTCTTTTGCCATGCCGAAGCTCTTTAGTT
TGATTAGATACTATTTGTCTATGTTGGCTTTTGTTGCCATTGTTTTGGTGTTTTAGTCATGAAGTCCTTGCCCATGC
CTGTGTCCTGAATGGTATTGCCTAGGTTTTCTTCTAGGGTTTTTATGGTTTTATGTCTAACATTTAAGTCTTTAATCCA
TCTTGAATTAATTTTTGTATAAGGTGTAAGGAAGGGATCCAGTT TCAGCTTTCTACATATGGCTAGCCAGTTTTCCCAG
CACCATTTATTTAAATAGGGAATCCTTTCCCCATTTCTTGTTTTTGTCAGGTTTGTCAAAGACCAGATGGTTGTAGATGT
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CAACGTGGGCTCTTTTTTGGTTCCATATGAACTTTAAAGTAGTTTTTTCCATTTCTGTGAAGAAAGTCATCGGTGGCTT
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GCCTCAATTTCAGAGCCTGTTATTGATCTATTCAGGAATTCAACTTCTTCCTGGTTTATTCTTGGGAGGGTGTATGTGT
CCAGGAATTTATCCATTTCTTCTAGATTTTCTAGTTTATTTGTGTAGAGGTGTTTATAGTATTCTCTGATGGTAGTTTG
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CTTGTTGGCTTAAAGTCTGTTTGTCAGAGACCAGGATTGCAACCCCTGCTTCTTTTTGCTTTCCATTTGCTTGGTAG
ATCTTCTTACATCCCTTTATTTTGAACCTATATGTTTCTCTGCATGTGAGATGGGTCTCCTGAATACAGCACACTGATG
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GTCGATGGTCTTTACAATTTGGCATGTTTTTGCAGTGGCTGGTACCAGTTGTTCCTTTCCATGTTTAGTGCTTCCCTCA
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ACTTACAAAGCTTAGTTTGGCTGGATATGAAATTCTGGGTTGAAAATTTTTTTCTTTAAGAATGTCGAATATTGGCCCC
ACTCTTTTCTGTCTTATAAGGTTTCTGCCGAGAGAGCTGCTGTTTGTCTGATGGGCTTCCCTTTGTGGACAACCCGAC
TCTTCTTGAGGAGTATCTTTGTGATGTTCTCTGTATTTCCTGAATTTGAATGTTGGTCTGCCTTGCTAGTTTGGGGAAG
GTAGATTTGGTCTTTTCACATAGTCCCATATTTCTTGGAGGCTTTGTTTGTTTCTTTTTACTCTTTTTTTCTCTAAACTT
CTTGTGCATGCATCACGAAGTTCTCGTGCCATGGTTTTCAGCTCCATCAGGTCATTTAAAGTCTTCTCTACACTGTTTA
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Fig. 6.45

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TTCTAATTTGCCATTCGTCCAACATTTTTTAAAGATTTTCAGCTTCCTTCAATGGGTTAGAACATGCCCCTTTAGCTTG
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TAGCTCGTGAGGAGCTGTGATCCTTTGGAAGAGAAGAGGCACTCTGGTTTTTAGAATTTTTAAATTTTCTGCACTGGTT
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GACAGATGTAGGCTGGATGATCTTGGGTGAGGTACACTAAATCTCAGTTCTCTCATCCGTGATACAATTGTACTCATCT
CACAGATTAAACAAGATAAATCTGTGGGGAGAATCCGCCCCAGAATTGAGAGAGGCTGTTCTCTGGGCACACTTGCTTT
ATGTGGTCTTTTCAATTGCTGTCAATCAGTTTTTAAAAATCTGGTTGCCCAAACACCATTGTGGCTATTACCATGACTAC
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Fig. 6.155

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TTCATGTTCTGGCTGTTCTGGCAGTACAAATGAGTCTGGACATCTACTTCAGGTAAATTGGATGTTTGATTAATTAATA
ACCATGTAGGCCATTTGTAGATGGAATCGCACAGTGTGATTAGTCATGGTAGCATGAAGACTTTAACAGGCCAATCCCT
ACATCTCAATGGCTTTATATATTACATATTTATTTCTTGTTCGTATCACATTACAATGTGGATTGGGGGTAAAGTGGGA
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TGCGTTCCTGAGTCTGTGTAGTCTTCCCATCTATTTCTCCTTTCGTTTCTTCCTCCTTCTCCTTCCTCCTTCCTCCTCT
CATCTTCAGAAAATGAAAGGCTCTGGAAGTCTATTTCTCCTTTGTGTGTGAGTACCTTTGGGACCAATGTAAAACATAGAGCAATAT
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TGCCTATAGGTTTGATGAGGTAATGTTATAAAGTATATTGCTCTCTAAAAATCCTGAGGGTTTTTGAGTGAATGATGTA
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TACATATATTCCATGACTTCAACTATATAATGTTGTAGTTAAAGTCATCATTTTAGAATCAGACTGAAGTGGATCTAAA
TCCTGGTATTATGACTTAATATTCACCTAGTCATTAGCAAATGACTCCACCTCTAGCCTAGCCTTAGTTTCCTTATCTA
TAAAAAGTATTTGATTAAAACCTACCTTAAGGATTTGTTGTAAGGGTTATAGGAAATAATGCTATAGGCTTTGGCAAAT
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CAGAAAAACTTAACAAAATGATTTAGTAAATAACTTAAGATCTTGGAGTGCCTCATAGTCATCATTGGAAACATAAGGA
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AAACATGCAGGGTTTGAACTAAGCATCGAAGAACGAGGATGATTTGAGAAAAAGTTCTGTCTCCTTGGGAATAATACAA
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Fig. 6.46

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AAGTGAATGTGTAAATATGTCTATAAAAACGATGCAGTGTTGGGTAACGGCAACAAAATTATTGGTCTCAGTCTCCCTT
GGCTGTTTTTTCCCCTAATTTCTTTGTGTTTTAGTTGATCTACTTGCAAGATGGAATAGTGGAGATAACATTTTCCTAG
TGTTCTTTTGAAGACACGTATAATTGTAGTCCTATTGTCCTGAGACTGAATGTACAGATGAACCCAGCCATGAATGGAG
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GGAATCTTTGGGAAGATAAAGATTGTGGTCAATATAACACCTATCTAGCCACCTTTTTTGCGGGTGGTTGGGTCAGTAG
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Fig. 6.47

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·GCCGTGCGCCTGCAGTCCCAGCTACTCAGAGGCCGAGGCAGGAGAATCGCTTGAACCTGGGAGGCGAAGGTTGCAGTGA
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CCTGCCGAGCCTCCTAAGACAGCACTGTGGAGTGGAAACACCTCTGATCTGGGAGCTGGATATTCTGGCCTCCAGACCT
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Fig. 6.48

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CAGGGTGCCTGATAATAANAGCATAAGGGACATACCATGAAGGAACAGGGGATCAACAATGACAGTTTTAGGACTGTTT
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GCGGATCACTGACGTGTGATCACTTTCACCTGTACATCATTATCAGTTCAAAGCCTCCCCCTGCCCCGTGGGCTTGCCG
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CATTCTTGTGTAGTGAGTATGGTTCTGAGGCAGTGGTCACTTTGTGANGCAGTCATGAAATAAATAGGAGAGAATCTGG
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CAACTAGTAACAGATCCGTAAAAGTGACAGCCACTTTTACCACCATCATTATAGCTCTGACCACTTTCCCTGACTTGAT
TTGGCCAATGTGTTGTGTGAGTGGTGGATGGAAGAAGCTAATACAGCTATGGCCTTCTTCAATAATGGTGAGGTACCAC
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<p align="center">Fig. 6.59</p>

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CTACCTGTTGTCTAAGGCCAAGCTGGCATAGATACCTAACTCTGGGTCTTCTCTGTCTGGGGGCAAAATCTCCAGCCTT
GAATAGGAGTCAGACCTGTTATGGTGGCTTTCACCTGAAGTCCCATCTACTCACGAGGCTGAGGCAGGCTCCCTTGAGG
CAGGATCCCTTGAGCCCAGGAGTTTGAGGCTGCAGTGAACTATGATCACTACTGCACTCTAGCCTCGACAACAGAGC
AAGACCCTGTCTCTAAAAAGAGAAAGAATAAAAAAAAGGAGTTAAAAGTTAATATAGAACCAGAGAGAGAGGGAACAG
GTCTTCTCAGATGGGGTAATCCCTGGTTTGACTTTCCTTCTAGCCCCATGAAGAAGACCCGTAAGAAAATAAGGGAA
GAAGCCAAAGAAATATGTGGCAGGGTCTTTCCCAGGTGGCAGCAGCCCAACCCAGTTCACCTTCCTGACTTCATAAAGG
AAGACACAGGAGGCTTTTCTTATTTTTACTCCTCTCATTAGACTCCACCATCTGCAAAATGTTGTCTAATAATAAACTC
TCAGGATCATCTATTTTAACATATCTGCTCCTTTGTGGGAAGAGTGAGTGTGAAAATTTGGGAACATGCGCAGTGGATA
AAGCTGTTATTTTATGCAAAACAGAAAAGAGCAGTGCCCCCTTTGGGTGGATGTAGCCCTGTACCAGTGCACACAGCTT
GGTATTCTCAAAGAAGCCCACAAAACAACCAAGGAGGCTGTTCCAGCTGGGCACTAGTAGCTGATTCTCTTGGTAACCG
TGGGCTGGGTTTTAGCTNTCACATGCCTGCTTGAGCGGAGTACCTTTGTGTGGTAAACAACCTGGACCACTCTACATGG
TCCCCTGGTGGTGGTAATGGGCAGGCTTAAGCGGTGTGATATTGGTTAGGGGGCTCAAACAACTCATGGAATGGGCACC
TAAGCAAGGTAGAGAGATGCAGAGCGGGGAGGGAGGGAGTTCTGAAGGGGAGTTTGGCTTGTTTGCTTTTTTTTTTTTC
TTCTATCTTGGCACCCTTTATTAACAGAACACAGTGTAGATTATGCATTGTATTTTGAAAAAGAAAGAGAAATAAAAGG
CGTAATCTTTGCTTTTGAGATGTTTTCAGTCTAACTCAGCAATTCCTTTCCCTCATCCTTTCTGCCTCACTACTTATTA
CATTTACTTCTAGTTCTTCCTCTTTAAATTGAAAAAGCAGAGGCAAAACCAGACACTGTGATAATGGAATTTTATGTTG
TGGTCTTTGATTACACTTCCCTTCCAAAGCTTCAATATTTCACATTGTTAATAATAATGTTCTTTGTAGGAAAAAATCC
TTAATTTGAAAATGTTGCAGATGGCCTTTTATCAGGAGCTATGGACTGTCAAAATAAATAGATATAATTATGAACCATT
GTTGTTGAATGGGCCATCTTAAATAATAGCAGTCAAGTTTCCAGAAATTAAAAAAAAAAACCTTACATTTTAAAAAGAC
CTCAGAATAATTAGTGGTCTCTGGTTGATTGGTTTTCAACAAGTTATTCAGCAGATGGAGGCTGTCAGATTTCTGCTGG
TACGATTATTTTGTAACTCAGTGGTGGAGGGGGGTGGTTTATGCAACTCAGCATCACCTCTTTGTTGAATTGGATTTG
GTTTAAGAGAATCCCACCGCACCTGTACATTAAGATGGAGCATGAGGAAGGTCTTGGGACCGAGCCATTTGAAGAAAT
CTAGGCTCNGGTGGGTCACTTTAGGAGTATTAGTATAGCCTTAGACATTAGCTCTTGGAACTACCCTGAAGGCGAAAAT
GTGAGACAGCTAATATTTCTCTGCAAGAATTCCTTGGGTTGCTGAGTTTTGGTTCTTGCCATCAAGAAAATGTTTGTAC
CGGAGCAAGATGTGGAAAACCCCAGATGAGAGGATATTAATGTAGATTCATGAGCTCTGCCAAAGTAATGTCATTACTG
CTGCTCCATCCCTGAAGAAAGATCTTAAACATATGTAAATAGACAAGACAAGTTATAAATTGTAATTTAGTATCTGGTA
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CTGCAGTAAATAAAATGTTATATTAGTGGAAAATACAAATGGATTCAGAAAATATAGACAAATCGATAGGCAATTGAAA
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GCTGCCACTATAATTTAAAATATATTGTTCTCCACTCTTTTTACATATTCTTGAAAGCAGTTCATTAACGGTGACCTTG
TGTAGGAAAAATTAGCATTTGTGCCAAAAAATTCTTTGTATGTGTTAGTTTGTGTGCATATTTTGGAGTCTTCATGTTA
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ACTCAGGAAAACATTCAATGAAACCTCTTTTTCAACACATAGATTAGTACCCTAAGAGAGTTAAATTAGTGTTTTACTG
TGAAACTATTTTCTTCCACACCAGAATCTCTATTGTATGTTGGCAAAGAAAAATGGACTAGCCATTCATTATCACCAAGAG
TTTATTACTTACTCTGGCCACCTCAGTGTTCAGTGGGTACCTAAAGTGAAAAGAGTTTTAAGTGCTACCTTTGCAGTAT
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TATCTTATTCCTTGAACTTCTTGATTAAATTGGTAGTGTTTTTTATTTGTAGTAGGCTACTTATAGAGTTCTTTTTCAT
AAAACTTGAAGGTTCTTCTTCATAAATCTTGCAGCTTCAAGGGGAAAAAAAGGAAAAAAGAAACTTATTTCCTCTTAGA
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Fig. 6. 50

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CCCTTACTGTCACACCTCAAGTATTGGACACAATGTGCGATGCTTAAAGACTCTCTTGGCTTAGAAAAGCCTTTCTCCT
CTCTTGGCAAATCAGGCAATGTGAAATCAGTAAGGGCCCAGTTCTCACTGTTTTCCTATGAAG  GTTCCGATGTGTGA
CACATCCACCTTATGGAGGTTTAAGTTGGTTGTGTCTTGAGGCTCACAAAAGCAGGTTGTGGAATTGGTATAAGCATTT
GATTTATTTGAACATATCTGGTGGTGTGTCATTGACTGACCCCTCTGTTATAAGCTTTCCTTCAAAAAAAGTCCATCAG
AGACTTGAAGTTTCACAATCAAAGCTTTGTTTACTGGGCATATTTTTCAGCCTCAAGGAAATCTTCCCTGGCTCCCTGT
AGAAGCAATAGTTACACCTTCCTTAGGGGAAGAACCCGTTTAAATGTGTTGCACATCTTCACCTCTTTTACTTGAATAA
GAAGTTTACAATCTTCCCTGTGGTATCCTTCTTAGGGGAACTCTAGGAAGAAATGTTTGATCATGAGAGTGGTAACTGG
AAGCATGAACTGGAGTACGGTGTTCCCCATTTAAAGAAATTTTGCTTAAATTTTTTTTTAAATGGCTTTCCAGAGGTCA
GGCAAAGGCCAGTTTCCCCTTGATATTGGGAATTTCAGGGTGGGATTAACAAAAGGCTAGAAAATGAAGATGGAAAGGG
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TTACTTCAACAAATATTCATTGAGTATCTCTGCCAGGCAAGAATCTGGGCACCAGAGTTACAAAAGTGAATGAGGCACA
TTTTTCTTCCTGAGGGAGCCTAGAAGGAAATATACAATTAACTACAGTCATTGTTTCTCACACTACCAAATGAAAGAGG
TGAGGCTTACCTGTTTCTCAATGTCGTTTTTCTGAGGACAAATAAATGAAGAAACTTCAACTAGAAAAAACAAAAATCA
TATTCACAGTTGGATTGTCTGTACAATATTGTCATCAGTAAATCATGTAATGGTATTATAGGACTTGGACTGAGTGTGA
CATTGCCAATTTCAGTTTGTAATTTGTTCAAACTTGCAAGTGATAATGTGGTACTTTATTTTGAGGAGCAAAACAATAA
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GAAAGTTCTATAAAAAGATTTCTCCACAATTCATTTTATAATAGTCCTGTCAGTAGGTTTTATTAACTCTACTTTTCTG
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TCCTATGGAACTGTGGCATGATAAACTTTTGTTGTTTTATGCCACTAATGGTAATTTGTTAAAGTAGCAGTAGGAAACT
AATCTGCGGGGTGGCTGTTAGAGCCTGATTTATAGCTATCTACCTTCAGAGCAGACAGTGTGACTGGGAGCCCCCTGTG
ATCACCATGTGGGGGATGTTATAGGTTAGATCAATCATTTATTCAATTATTTATTAATAAACATTTATTGCACATCTACT
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GCGCTAAACTCAACCAGAGGTCACAGAGTACTGTTGGCAAAGGTTGGCCTCTTTTTCTTGCTGCACGTGGCTCGTATTT
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GTTGGACACCCAGAGCTAAGAAGTAGAGAGATGTAGGTAGAAGGGCCAGCCAATTGGCAGCAGTAGGCTGCAACAGCCA
CACACTTGCCCCGGGAAGTGGGAAATAGGAGGGATGCTAGAGCTTGGTTTCTAACATGGCAAAGATCTATGAGAGCGAG
AGAGAATAGTAATGGAGTAAACCAAAGGAAGAAATAAGATGCCTCCAGAGGGATATGGCAGCTTTAAAAACATGGCCTCA
AAATCCTTTGACACTCCTTCCATCAAGAAATGGGGTCTATGTCTCTTCCCCTTGTATCCTCTTGTGACTGCTTGACCAA
TGGGATGGGGTAGAGGTGTCAGTTTCTGGTGCAGACCTTCAGAAATGGGTCGCTTCTACTTTGTGTATCCTGAGAGGCT
TGCTCCCAGAACCCAGCCAACATGTTGTGAAGAAGCCCATAGAAAGGCCCGTATGGAGAGGAAACTGCAGTCAGCCATG
TGTGAGAGTCATCTTTGAAGCAGATCCTTCAGCCTGTCAAGCAATTCCAGCCGACATTACATGGCACAGAGATGAGCTG
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GGTAGCTTGTTATATGTGAACGATTGTAACAGGGTGCATTTTAGGTGACTAGGAAAGAGAAATACCAGGTTTCATGCAA
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AAGCTTAGGGAAGGAAGAGGGAGAAGCACTGGACTTTCTATTGTTAGACTTGTATTTTATTCCAATCCTTTCATAGAT
TTGCTCTTTGATTGGAGATAAAGTGTAAAACCTCTCTATATCATATTCTATATCTATATGATATTCTATATCATAATTT
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TTGATAATGGATTACTTAATTAGGTCGCTGTGGTGTTATGATATAGATACTTGTTTTCATCCATGGTTCCTGGCTTCTA
GCTCCCATAATCCTTGTTATAATGTTGGGCACTTTAGGCCTCAGGAAACGGAATCTTTCCCTCTAACCTTCTCCTGTC
CTCCTTTCACTTGCCCAAGGCAGGACTCTAATCTGATTGTCGGTCAAAATACCCTCATTCCAGATCCTGTCCTATGCAG
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Fig. 6. 51

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GCACATGGATGAAGCTGGAAGCCATCATCCTCAGC..ACTAACACAGAAACAGAAAACCAAGCACCGCATGTTCTCACT
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CCCATTTTTTTTTTAGAAAAAATAAAGAAAAAAACCCCAAAAAACCAAAATACCCTCATTCCAGAAAGAGTCCTGCTCTA
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GCATCTTCCCTCATACCTCACCCTACACATCTGTATCCTTTGTAATATACTTTATAATAAACTGGTAAGGGTAAAAGTG
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GGGGGCAAAACTCCACACTTCGGGTAACAGAAGGCTTCTTCTGTGTTGATGACTGTTGTTGTGGTGGCGTGAGAGTAGA
GGAAAAACACGGTTTGAGAGAGCTTTTCCTGACACAGAAGCCCAGACCTTTAAGGGCAAAGTTTTGTCCATAAAACAAC
CCTAGAACAAGGTCCCCAGGGAAAAAGCACCAAAGGGTATAGGAGTGGAAAGAGGGGGTCAGTACAGGCATCTCTGCCC
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CGCAAAGATTGGTATCAGTTTACACCAAAACAGAACACACTGCTGGTGAGCACAGGGAGTTGGAAACAATATACACAAA
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TCTTATGAGGTCAGCAAAACATTTCAAATAGCAACTCCTTGGCTGCCTGCAGTGGGTCTGCAAATGTGAAATAAAACAT
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TTACTGCATTGGTAGATTTTTATCATCTCAGATGAAAGAATCCTAGTGTTGGCCAGGGCAAATCCAAAACACAGGTGAT
GAGAGGCAACACCCAATGTTAAAACCTGATCCACTGTGGTTATTTTTCTTTCCCATTGGGAAACTATAAGGATGCAAAT
GGGCAGGAGAGTAAAAAGCAGCAATCACGTCTGCATGGACTGAAGGCAATTTAGTTTCTATCAGACATGGTGACAGTGAT
CAATGCATCACAAAATCACAAACAACAGATGCCAAGCACAAACTGTGTCACAGATCCCAGATCAAAGATATCTACCATA
TGTTTAGCTTTCACATCAATGTACATCAAGTCAGTTTTGAACTAGAAGCCTTTAGATATACAGTTCATTACTTCTTTCT
CCTTTTTCATCCTCTTTTGTCTCCAGGAAGTTCCCAGAATCCGAATTTAATCATTATTCAGGTTTGGGTATATTTGCAT
ATAAATGCATTGCTTGTGATGGATCCAGTGAAAGTTCATGCGCAGGGCTGCCCCTGCCCAATGCTGCACTGGCTTTCTG
CCCCATCAGTGCTGGAGCTGGCAGAACCCCCCTGCAGAGGAACGGAAGCAAGTGACCTAGAGGGGCCTGCAAAGAGTCT
TAGGAGGGCAATTACAACACGTGTTTCTTCCTTTTAGGAGTATCTAATTGGGTTCAATGTGAAGGCAGCTCACTGTCAG
GCACATAGAAGAGACTCTGAGTTAGAGAGGTTGAAATGTGGTCACAAAATATCCATGAATGTATTAAATCCATAAGGAT
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CAAGGAATTGAGGGAAGGCTCTTGAGAGAGAGACAGAAGGCAGGGCAGTGGCACAGGTGGAAGTGCCCAGCTAGGGTA
AGATGAAAATAGGATTGGGAGACAGAGGTGGACATTCTGAGAAGGGGGAGTAGTAAGTACAAAACAGAGGTGGCAATAG
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GCTGGAGAAGAAAGTAGGACACAGGTCATGGAATGACTTGCAGGCTGCACCAGAGGCACTTGGATTTTATGAGAGCCTT
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TCCCATACTTGCCATTCTTCCATACCCTCCTACATTTTGCTTCCTGGAATCCCTTCTCGAAAAACTACTCATGCTTAAA
TTCTTAGCTCAGGCTCTACTTCTGGTGGAATCCAATCTGGGACAGGTTTTTGGCTTGTGGAGGAACAGTGGCAATGAAG
GCGGAGTCTGGATGATTTTCATGAGAATGCAAGAGAGAGTGGAAGACTCTGGTTCAATTTAGAAAGCACAGTTCGTTGC
AATTATGGCCAAGCTGTATGTATTCTGGGAGACAGGGAAGGACTTTTCGATGATTCAAGGTTTAGGTTTGTTGACTGGT
GGGATCCATTAAGCAAAATAAGGAATACAAGAAGAGGAATAGGATTTTTGGTGAAATGTGTGGGATTTGGATTTGATTA
TGCTGAATTTGAGGTATTTCCATGGACAGTGTAGGCATTGGGGAGAGGACTAGTTGCGACACCCCAGTTTGGAAGCTG
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GGAGTGATCTGTAGCACCAGGCAAGGGCCAGGGAGTGATGTCGTAGAAGCCAAGAAAGGAAATCATGTCAGGAGGAGA
GAGGAAGCAAACAGAGTCAAAGACTGCAGAGAGGCTTGGCAAGGTCAGACTTCAGGCTCCCATTGGATTTGGAAGTTGA
GTGGTCATTGTTTATTTAAGAAGTTCATACTAATTTGGATCAATTAGAAAAGTAGTAGAGATTACCTCCACTTAAGACA
TTTAGTTTTATTACTTGAAAACTAGGCTAACCAATAATTGCCTAGGACGGGATTGCTTTAATGAATAGGTAAGAATAA
TCAGTACTTGGAAAGCTTCAAGTTTGTTCACCTAATTAGGTAAGGGTTTTTCTTTTAAGCTAGTTGCTGTTTTTTAGTT
CTGATGCCCCTTTGAATTTTGAATAACAATTCTATTTTCTTTATACTATTTAGTTATTTGTGGTTAAGAATCTGGTCTG
TAGAGTTAGACTGCTTAGCTATTACAGTAGCTTCAACACTTACTCTTTAAGATCCCTGGGTAGGTACTGTGCTTCTCTG
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Fig. 6.'52
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TGCCTTGATTTCCTCATCTGTAAAATGGAGAGAATACAACTTTTTTTTCAGAAGTTAATTCTTAGTACCAAATGAGCTAA
TTCATATAATGTATTTAAAGGCATTTTAAATGGCACAGAGTAAATGATCAGCACATTTTAGCGTTAGAAATATTTGTTA
GTATTTTTTCTATTTATAATTTGTTACTATTACTAAAAACTGGGAGGCAGAATAGTATAAAGTGATAATGATTGAACTA
TAAAACAGATTAGGGTTCTGCTCTCAAATTTTCCGCCAACTCTCTGAACTGGGACAAGTCCATTCACCTCTCTGAGTTC
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TAGTGCATTCATTTCATTTGCTAATGTTATAATCTGTACAAAACAGCCCACCTAATTATTCTTTTAGATGTTAAACATG
ACACATGTAAATAAAATGATAATTATAGAATGTGGTGTTTTCTTGTATCTTACATTTTTTAGATCTGAAAATTGGTCCC
TGTTCTTATTCTGCATGTACTCCAGTGAAACTTTCCCTTGATGAGTTATTTTTCATGCGCACACATGGGGGAGCTTTTG
AGACACTAGTTTGAATGTTACACTTTGAAGACTTTCTCAACAACTTGACCCTAAGATGATGGACTGGAACTCTTCATTG
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CGGTGAAATGGGGGAAGTCAGCCAGAAACTCTGATTTTGACAGCAACATTATTTAAAGGAGTTTCTGAAATGATCAAAC
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GGTAGCTAAAACCGCAAATAAACATATAGAAATAGACCTGGTGTTAGCCACATAGATACATGAAGTAGCATCAGGAATG
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TAAATACAGCTGATTTTAAAATGTTATTTAGTATCAAGAAAAAACTCTTTAGCAAATATTTCCTGTGAAAAGTATACAG
TATAACTAGCTTAATATATAAAATTAAACAGAATCTTTCATGATGAATTGAAAATAGGTAAAATTTGTTTTCCTTCTCA
TTGTTTTTTCATATGTAGACAGCCAAGTTTCATCCTTCACAGTTTGTGTTCAACTCCTTGCTGTTTTTTTCTCTAATCT
TTTCCTCAGTGCCTTATTCATTCTCATGGTTTTAACCATTAACTATATTCCCATTCTGGGCTTTCATCACAGATTCCAG
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TCCATCCTCCACTTGCTTCCTCGGCTTCTTTCTTTACCCACCCTCCCAGCTCACCAGTTCCTTCTGACTTTCCCGTATC
TGTTCACTAAACTACCATTTTCTGAATCACCTGGCCTTGGAAATTTGGTGTCATCTTTGACATTAACCTGTCTCATGTA
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CACAATTATGTTTTACTTAGTGAAAGAATACAGACTAAAATCTGAAAAGCAAAAGATATGTGGGGAAAAGTCCAGGAGA
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AGTGTTTCAAAGAAATCCAGGTGCTATTCAATGACAGTGAGACGTCAGGCCAAGGGGGTGAAGGGAGGCTGAAACCCAG
TCCATGCTTTGCTCACAAGCCAAGTCCTGAGATGGGATGAGGAGAAAGAGGTGTCTTTTTCTAATCTCATTAAAGCACT
GAGTAGTGTGGTGGGGGCTTTTCTTCTGGGGTTTTGTTTACTAAAAGACTTCCTACAAAGAACCTGTAGGCCCCACAAA
GATCATATGCATGGACATTATTGTAGGGCAGCAGGAGAAAAATGCTATTTTGGTTCTGCTTTCTAGAAATTTTCAAGTG
CTGGGCTACCAAGTCAACTAGCTCCTCTGCATCCTTTAGATGTCTGTGTGGCTGAGGACAGCTTCATGAGATTGGGTCCTC
AGAGCTGCTTTGCACTTCCCAAGAATAGACCTGTGGACCATGTCCTTTTTGTCCACCCAAGTTTTATTTATTTTTGGGA
CAGCACCTCTTTTACCAGAGAAAGTAACTCTTGCGGCTAAAAATATACCGGAAATAAGAATGAAGAAAAGTAACTGGAT
AGCAAATAGCCAAAACGACTTCTTTGGTTGATTGGTTTTAATGTTTTTTTTTTTTTTTCTTTTCCTGGTTACAGTTTTAA
```

Fig. 6.53

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AAACTGCTCCACAAGAAATTTCTTCCAAATTCAGTGCCAATAGGGAGCATTGCCTATTAGCAGGCACACAATAGTTAAG
AGCAGAAAGGCATGCAATAGGGCTTTGATTGTCCAAGGCAGGGTAACCATGTATGTAGGACCTTTATTGCTGCCTGGGA
GGTGCTTCAGTGTGGGGAGTAAATGCTCAGTCAAATTTGGCATAACAGCTTTAAAGCAAGTGGAACCACTGTGCATGTT
GTCAGCAACATACTGCTTTCTTTCTTTCTTTCTTTCTCTCTCTCTCTCTCTCTCTTTCTTTCTTTCTTTTTCTTTTTAGTTTT
TGAGTCGGAGTGTCGCTCTGTCGCCCAGGCTGGAGTGCAGTGGTGCTCTCTTTCTTTCTTTCTTTTTCTTTTTAGTTTT
TTCAAGTGATTCTCCTGCCTCAGCCTCCTGAGTAGCTGGGATTACAGATGGGCACCACCATGCCCTGGCTAATTTTTGT
ATTTTTAGTAGAGATGGGGTTTCACCATGTTGGTCAGGCTGGTCTTGAACTCCTGACATCGTGATCCACCCACCTTGGC
CTCCCAAAGTGCTGGGATTACAGGTATGAGCCACCATGCCTGGCCCTTTTTTTTTCTTTTAAAAGTATTTTACTTATGTT
TCTATTTTGTGTCAAGGGGGCTGGTTCCTAGATTACTGAGGGAATAAGCCAAACAGAGCAAAATTTGTGTCACTCATAT
TAAGGTATTAATTTCAGCACTAATGAGATGTTGAAAAACATGGCAAAGGTGATTAGAGGATATTGAACGAAGGTAAATT
AAGGGTTTTAAATTATAGAGAAGATGAAGTTTGATCAACCCAACATGTGTCCCCTTTACTAGTGGGAAGACACAAGGGC
ATTCATGTCACCATGAAATATTCTTATTAATTTATTGAAATACCAGTGTATTTGTGGCATTTTGGGATTACTATCATTA
CCATAACAGAACTGTTCTTGCTTCATATTTATGCGCTCCAAGATTGCTCTAAAATTCCTTCATATCCAACTGGGCAATT
ATGATTCACCCATTGGACAAAACAATATAGTTCTATTCTAGCACTCAAAGGATGTAGTCATCACTTCCAAGAGTGAGCA
GACTTTCTGGAGCCAGGTCAGAAAGAGACATTAATGTTGTCTTCATTTAAATACTATGTGTAGAAAAGACCTATGTATT
TCCCATTGAGAAGATGGATCATCTTTTTCAGGAATTGTTTTTCATCTGTAGTCTCATCAGATCCTGTTAATATATTTTA
CTTTGCCTTTGCTTTTTTTTTCCATTTGAATATATCTTTTTCTCCTTTCAGCATGATGAAACGTCATTTAATTATAAATT
TTTGTCTTCAAGACCTCCCTTTAAGCCTTTAACGGGGAGTAATATAATGACTACTGGTAGGTACAGTCAGTCAGTCAGT
TACCTCTGGGATCATTCTTTGCAGTTCTTGGCTCTTGGTTTCTGTAGGAGTTGGCAGGTAGATGCCAAATTTTCTGTCT
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TGGCTAATAAACAATATTTGGTCAGAATTTGAAGCTGATAGCTTTCTCTCAATGTTCTAAATAGATTCTATTCCTTTAA
AGTGGTTAAAGTAGAACAATCCTTCAATCCCTTGATCTGATTTATGGCAGTAGTAATAAAGAGGATTGTTACCCCAAAT
GGGCCAGTTGTTTATGCTTGTTTATTCATTTATTCTGGAATGTTTGAGTGTCCACGATATGCTGACTCTGTGCTAGATA
CTGGGGGTTCAGTAGTGAGGTCCAAGTCAGATATAGTCTTTGTCCTCATAGTGTGTAAAATAGTGAAATCTAATGGGGA
ATACAGAAATAGGCAGTGACAATAAACTGTATTGTGAGGAATGCTATAGGCGAGAAGCACAGTGAGCTCACAGGAGGCA
CATGGACAATATTTGGTTGCCTGTGTTTACCAGGAAGTGTGCTAAAAGCTGAGTTTACAGTTAACTAACAAAATAGACA
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GAGCCTATTCAACCACAGCAAGTGTCTTGACTTTACCTGGAGGGGAATGGAAGAGCTACTGAGTGTTTTAAACATGAA
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CCTGGAGACTAGGGAGTAGGAGGCTGTTACAGGAACCCCGGCTAAACAGTTCACTATAGAGTAATGCTGGGAGTGTGAG
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AGTAGCTGGGACTACAGGCGCGTACCACCACACCCAGCTAATTGTTGTATTTTTAGTAGAGATGAGGTTTCACCATGTT
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CACCATGCCCAGCTGGTAGTATATTATAAAAGAAATGATATTAGATACACTGTATTTTGTGTAGCCTGTGCCCTCCAAA
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TTATCATCTTTTTTTCCCTTTTGAACTGGGAACAATGCCCTGATGATCTATTTGGAGAAGAGATATGATAGTATCTTACG
TGACCTGTGCTTGCTCTATTCTGTTTGCTGTTTCCTGTTTAATTCTGATGGTGTTGTGGATAAGTGAGATGACCTTCCC
TTACTCTCAGACCGAATTATTCTTCGGAGCTTTGCTGAGGGCTTAGAGTACCCTAGGCAATATAAACGTTTCTTTGATA
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Fig. 6 /54

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TGCTCCTCAGTATTTCTTCTTTTTAAAGAATCCCCAAGAGGATTTGCACCTGAAACTGTAGCTAGTGCCATTCTTTTTT
GCTCTCCTTTTTCTCCTCCCTTCTTTCTAACAATTCAAATGTTGATGGTGTTTTTTTGGGGGGAGAGAGATGGCAAAA
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TGATAATGTGAACAGCTAAGTTCCCTGGGTGGCTTTAAAGGGAAAGGGACAATTCACTCATTTTGTGTATTTACTAAAA
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AGAGACCTTCTGATTGAACATGCAAGTTGGGGCCTTTTAAGAACGAAAGGGGGTGCTATGAACACCAGGGCAATGGCTG
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AGGCTCTGTCCTGTTAGGGGAGCTGCTTTCTGGCCTAGCCACTCTGGACTGAGTAAATTCACAGGTGTTCACTCAGCTA
GGTTTTCTCTCTGTGATAAATCCTGCAGGGGTTCTCTAATTTAGGACTTTACAGAGGCACTAATTGAAAAAAAATGAAA
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TTACCAGTTCTTTCTTACTAAGAGTCTGGTTAAAAACTCATGAATGAATATCACTCACAGAATGACCTCGATTCACGTC
TGGTCTGATTTTTTTTGTGTGTGCATGTTGACTTCTCATGTGAGCTTTGTTCCTGCAGACCCCACACTTAGTCTTATA
TGGTCTTCGTCCTACTCTGCCTTGTTCTGGGGTGTGGCTTTCATATGTTGGGGTCTTATGAATAGAAGCAGACAAGCCT
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Fig. 6.55

CAATCTCATTAAACACAAGGTTTTTGTTGGTTTTTCTAAAGGCAAGGTAAGGAAGTCCCTATGACATGGACCATGGTGC
GCATGCTCCTCGTGGCTTGTCCCTGGGCCTGGCTCATGAGGTTTCACTCCCACTCAACCCC TCTGCATCCCGCCTCT
CTCTGCCTTGGTGCTCTGGTCAGTTGACTCACTGTTCTGTCCTGTGCACTCTGTTCCTAGCTGTGGGAGTTATTGGGTA
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CTGAACTCACCTCTTCAGTCACAAACAAAGCAGCAGCACCTGCCCTGGTGGCTTCAAACAAGGAGGACAGAAAACATC
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TTCTGAGAAGTCTATCACGAGAACTGCAAGGGGGACATCCCGCCCCCATGATGCAATCACATTCCACCAGGCCCCTCCT
CTAACACTGGGGATTACAATTCAACATGAGATTTGGGTGGGGA_TAGAGCCAAACCGTATCAGAATTTTCTCAAGGAT
CAAATAATTCCCTCTTCTGGCCCTCACGGAAAATGAATGAATGCTAGAGCAAAGGAAAGCACTGCTTTCTCAATACCAA
CAGGTGTTGATGGTAATTCCTGTTTCATGCATTAAAGCAAAATGTGGCCAGAGGGAGAAAAGGGTCCCCCTAACCCCCA
CATCCTGCCAACCTCCCTGCCCCAGAACACAGGTCACGATATTCCCAGTGAAACTTAATATCGTGACTGCCAGAATGTT
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GACTTTTCCTATTTTAAATAAAGAAAACTTATCGTTGTTATTTTATAAAATGGGTAAGTTTATGAGCATAATGGAGATC
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TGACAAGCATACTGAGCCAGGGCATCAGGACTCCCACTGGAAGGCCCAGAGTAGCCAGGTCCCAGCAGGAGGGGAGAAAC
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TGTGTCAGTTTGTGACCTCTGCTCTGGTATATAAGTTTGCCCTTAGAAATATGTTCCTGTATTATTTTAAAGTTGAGTG
GGAAGCTTCTCTACTTTTAATGAGTTTTAAAGGAGAGATGTCATCTACTTATCCCTCTATCATTTGGAGGATAGGCTAG
TTATGTCACTTGGATTATACAAATAAACATTGTTGTGGGTTATTTAAAAAAAATTTATTTTCTGAAGCGAGTCTATCAA

Fig. 6.56.

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GGTTTTTAAAATGTGGTTCTTCATGGACTTCCAAGGAGGTTTAGGATTGGCAGTGAAAAGAAAGAGAAGAATGGGAAGG
ACACTTACATTTCTGCTTCTTGGGATAGAGGTAATCTGTATATTTTGGTTTTAAAAACAGATATTTAATCTTTTAGGCA
CATTAATCTACATTTCTAAGAAGAAATAGGAAACTTCTTGATAGAAATGCCACCTGTTTTAACGACTAAGCACCAATGG
CACTCTGCCAAGTTGAAAACAAATCTAGGTAAATTTAAATTGGGCTGGCATTTCACAGAACTGGTGCAAGCATGGTTCA
TAAAGTTTATTTTTTTGCTTGAGAAAAAAATGACATTTTACTGATTCTGTGTAAGAAAACCTTTAAGTTTTTATCAAAG
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TTAATACAAATAACTTAATATTTTACAATAGAGTTGATTATTTTTCTAATGCTTTCCTAAAGAAATTCTGTAAAGATCC
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TTGACAATGATCTTCTGATAGATCCTGGCTTGGGTAATACATTACACTGGCAGAGTAAGAATTCAGAAAATAGCTCAAC
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TGTTTCAAACTCAGGAAGGCCTGGGTGAATTCCTGGTAGAGCTGGTAGAACTAGCTTTCTATTATGAATATTGAGAAGT
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GTAACTTCCACCTACTTGGGCCAGCCCATGTAGTTCAAGGTAAAGATCTTGGATTTTATTCCACATAAGGTAAAGATCT
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AAGTACGGCCTTAAAACAGAAAAAGAAGCAAATATATACTACTTGAAAGTCAGACTGTATGATAATCTAATGGTAGATA
TAGTTATATAAATCATACACACTCAAAGCTTTACCTCTATAATTCTGATAAGAATGGGAAGGCTGATGACATTTTTGCA
GACCATTAAATAAATGTAGTGAACTTCTAGTTTGCTTTGAGCCTATCTGCATGGCTGATAAATGCTTTTCCAATAGAAA
GAAGGGAAAACATGCATTCAAGGTGATAAGCAACCGGTGATAAGCACAGCTAGAGACAGTTTTTAAACCCTGAAACTCT
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AAATAAAATAATATTTCTTTTTTTTTTTTTTTTTCTTGAGATGGAGTCTTGTTCTTCACCCAGGGGGCTGGAGTGCAGTGG
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CTCAAACTCCTGACCTTAGGTGATCCACCCACCTCAGTCTCCCAAAGTGCTGGGATTACAGGCGTGAGCCACTGTGCCT
AGCCAGAAATAAAATAATATTTCTTATCTGTACCCTAAGACATGTTGGACAGGAGATGTCCCATGAAGCTGATGAAATT
TTAACTAATGCAGTAAATAATCCATTGAGATTTAATCCTCATCAATCATCATCCTTATTGTTACCATGTGGCAGAAAAA
TGTGAAATTCTGTTAAATGCTTCATCTTGATGGAAAAATATGAAATCCTTGTAAATAAAGCTTGAAATTCTCAGGTTCC
CAAAAGATGACTCATCTGACCTTCTACTCATCTTTTTAAGCACATGAAACAAAAGAAAGCAAATGGGTTTCTGTGGCCAA
AAGAATTTCATCAACACCAGTCCATGGGAACAGATCAGACATAGGGAAGGTGGGTCTCATGACACTTTTTGGTTTATTT
CACATCTCTTGAGCAGAGCTCTCAGTATAATCTCCTCAACAAGAATGTGTATTGCATTTCTGTAGGACATAAATTGCCT
TCTCTCCTCACTATTCCCAGCACTGATTACAAGCTTCAGCAGAAAGCAGAGTTTTAAATTCTTGTGGAATTAATAAATA
GATAAGAGTAAAGGTATTTAGGGGGAGACAAGGACATAGCCTGTAATTTAGGTGAGCAAAATCAGTAACTGTGAGTCTG
GGCTGTGACATGGTCCTGGCTCACTATTATTTTTTTACAAGTTTTAGGATCTTGGCGTTTTATTTGCTTTATCAATTAC
TGTGGACTAAGATGTGCGACTGTGATGTGGATTAAACAACAACAATTTATTTCTTGCTCATGCTGTCTGTCCATTGTGG
GTCAGCTGAGTGCTCTGTTCATTCTTGTCACTCAGGCATGGGTTGATGGAGCATCCACCATCTCAAACGTTGTTAATTA
CCATGCTTGGGAAGAAAGGAAACTCTAAAGGATGTTACCCCAGGTGGTTAAATAAGCTCATGTAGAAATGGAATGTGAC
ACTTCAACTCAAAACTCATTGAACAGAACTCAGTGTGCCTGGCCCCACCAAAACCATCCACAAGGGAGGGCAGAAAGTC
TAATTGTGAACAGAACTAATGATATTTACTTAGTATTTGTCTGCTTGATTTTTGGCACAGACTGATTTATTAGAGTGGA
GAAGGTAAGCTGATTATCAAAGAATGTCTTATTTTCCTCATGTAGTTACCATTGATTTATTGAATCATGATTTCATTCT
TTAATATATTTTGCATCAATGATTGATTAATTTTCACAGCTACATCAAGTAACTAATTATTGATTTACTAATCAATTAT
TAAATTCTACAAATTAAATAATAATTACAAACTAAATTCAATCCTTGAATGTATCCAACATTTATTGGGAACCAGTCCAT
GGGAACAGATCAGACATAGGGAAGGTGGGTCTCATGTGCTTGATCATCTACTATGTGCTTGATCACTATGCTAGATACT
ACATTGTATGGCTTTGCTGAATTATTGTTAAGTCTTCACTCACTTAGCTGAGCTCAGAAATGAATACATCTCAATGCAT
GCAGAGAAGGTTCTTATTGAGGCTTACTCACTTTGTAGGGTGTTGGTGGCGTTTGAGATTGTGTTTTCAAAGTGCTTAG
CACTGGCGATAGTGATGTTCATAATGATGTTGCAAGTTGACTAGGATTTGATAGTTCATTTCCTAAAATAATTTTTATT
CAACTAATAATTCAAGTAAACACTGCTCACTTGATGATTAGTCGTGGAGTGAAGACTGCGTTGTGAAGCTCACCATTCA
TTCTCTAACAGTGGTCTCCTGAGTTGGGACACTTACATAGGATTTATTTCTGATGATAACAAGAGCACAGATGCCAACC
ATGCATGTGATAGTTTCTGGGGAAGACTTGATGATTCAGCACTGTCCCTATAAAATGACAAAAGAAGACCCACATATA
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GCTTTGAAGTTGGCCTCTTCGGTCTCCAGCCACAGAAGGGATTTTCTTCGCTGACCACAGTTCCCCACGTTTTCCCTTC
AGTAAATAGCAGATGTGTGTGGTGGTAAGCTGGTTTTCTGCCATTGCTGTGTGAAAAGGCAGATATTCTGAATGAGGAT
TGTAAGTTATTTTTGTGTTACACATTCTTCTTTAGCTTTTTTTCAAAAGCAATTGCATTTTAGTTGAATGTGAAAATTTA
CTCTTGATGATTTTCTCAGATTAATGCTGTCATCATGTTAATGGGGTAATGCTTCCTTCTGCAAAGATCTTGGAATTTA
GGGCTTCTGGTCAGTGTTTTACAGCTTTGTCTGAGAAAGAATGTAATATGAGTATGTTTTTGGTACATGACTTCAAAAA
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Fig. 6.57

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GCCAAGTCTGCTTTGCCTGATTTTTGTGTGAGCAATCCCACCTGTCTGATCTCACCCCTGCTCCCAGGTCACAGATAGA
GGAGACCTTGTGCCCTAGCAACAAATCACATCCACTTTAGCTCTATGGGGACATTTCAACAACAAGGTTCTTGTTGNGG
CTTACCCACTTTGTAGGGTATTGGCGGTGTTCGAGATTGTGTTTTCAAAGTGCTTAGCACTGGCACAGTGCTGCAACTT
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GTTGGGTAAGAAAATATGGCACTAGAAACATGGGAGAAGGNGTGTGTTTTGTGGGGGTAGATGGATAAGAAAGAGAACA
GNTGGATCAGACACCAATTTACCAAATGATTCCATAATATTAGTGCACAGATAATGCACAGATAGTGTGCAACAACAGC
TTTGGGAAAGGCAAACTGACTGTCTATAAAGGTCAGAAGACATTTTATAAGATGTGTTTTACTGAAAGTGTTGTCCTTG
TGAAGAGGTTTTGAGCAAATATTTTAAGAAGTCTTTAGGGAACAAGCAATTTTCCTTTTTGTTGCCCTTTCTAAGAATA
AGACATAAATAGGAAGTTCCATCCTTTTATTTGTTCTATTCAAATATTTATAGGGGAAAATCGGCTACTACTTCTTTAT
AAGACCCACCTTTGAAACTCATTATTTTTCCAAGGTGCCTTGTTCTTATAACAATATTTTCTTCTATGTTTGTTTATTT
ATTTATTTTATTGAGATGGGGTCTCACTCTGACACCTAGGCTGGAGTGCAGTGGTGCGATCCCAGCTCACTGTAGCCTC
TACCTCCGGGGCTCAAGTGATCCTCCCACCTCAGCTTCCTGAGTAGCTGGGATCACAGGCTTGTGCCACCATGCCCGGC
CAATTTTTTGTGTTTTTGGTGGAGATGGGGGTTTTGTCATGTTGCCCAGGCTGGTCTCGAACTCCTGAACTCTAGCAATC
AGCCCTCCTTGGCCTCTCAAAGTGCTGGGANTACAGGCGTGAGCCACAGTGCCCTGCCTATAACAATATAACAATATTC
TGTTTAGGCTATGGAAGACCACATATATTCTTACTTAAGCACTTAGAATGGAAGCCACCTGAAGACAGAGGTTATATCT
TCACCCTTGTCACCCACACATAATAAGTGCCCAATTGATGTCTTTTGATTAATTAAATGAATTAATTAGTGAATGAAGA
CAGACCTAGGACTCCTGATAGAAAAGAATAAAAACTTGCTTCTGTTTGTCCTTCTACAGTGAAAAGATTTCCTAAATA
TCACATATAATACCAGCACACATATCTTGTTTGCTAGTGATTTGATTTCATTTATGTCTGTTATTTAAATTCAGGGAAA
AACATAGGTAGTTTTTTCTTGTAATATCTCAACACTGAACTTGAATGAAACCTTATTAAATAATGCGCAAAATTAACTA
TTATATTTTGCTGCTAAGGGGCTTTGGCATTTTGCTTTGGTAAATTATGTCGCTGGTCTGGAAGACTCAGGCCCTGGGA
TGACAGGCCTATGTCACAATGGGTGACATGGGTTGGGAAAGATGGCTGGCTGTCCTGTAAAGTGCCTTATGAGTGTGAA
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ACATGTCTGATACAGTCAGCCAAAACAAGATTATTTATCAATGCCTCTGCCAGCTACCCAAGATGGTTAAGAGAACCAA
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AAGGGCTACTCCCTTACTGCACCTGAGTCCTCACATTCTCAACCACATCAGTCACTAGGAGCGGTCAAGGGAACACCA
CAGTATAGCAGAGTGGTCAAATCTGCAGGCTCTGTTGCCAGACTGCCTGGGTTCCCCTTCCTTTTCTATCACTTATTAC
CTTGATAGTTACTAAACCCTCAGTGCCACAGGGTTTTGATCTATAAAATAGAGAAAGTAATAAAAATGTTATCTCATAG
GGTTGTAGTGAAGATTAAATGAGATACTGTGTACTTACTATGTATTGTATGGCTTTTGTTAATAAAGGCAGGGTCAGGA
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CTGGTACCCTCTCCTTGCCCTTTCTGTCTGCTTTCCACTGGGATTGAAATGCTTTTCTTTCCTCCACCTCTGTTTCTAG
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TCTTCCTCCTGAGCCTACTCAATGCAAAGATGATGAGGATGAAGACCTTTTGAATGATATGGTTCCACTTAATATATAG
TAAATATATTTTCTTTTCCTTATGATTTTTCTTAATAACATTTTCTTTCCTCTAGCTTACTTTATTGTAAGAATACAGTA
TATAATATATATATAATAACAAAATACGTGTTAATTGGCTGTTTATAGTATTGCTCAACAGTAGCCTATTAGTAGTTA
AGTTTTTTGGGAGTCAAAAGTTATACTCAAATTTTTGACTACACGTGGGTTAATGCCCCCAACCCCCACCTTGTTCAAG
GGTCAGCTGTAGTAGATTAAAAGGTTTTTTCATGTCTGGAAAGATTCTCTATTCCTGGACTGGTTCAAGGAACTTAGAGG
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Fig. 6.58

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TCAATCTGAATTCCAAGGAGTGAAGGTACTAATGGTTTGATATCCAGGTGTACATGGCCCTCTGCCAGGAGACTGTAGG
GAAAGTAGAGATTATCATCATCCAACTTGTCCAGTTGTGCAGCCAGATATTCTTGCTGACAATGTGAATATTTACCTATGA
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GCAGTAGTGGCACATAATTTCTTGATCTTGTGTGTTTCTGTTACGGTAAAAATAAAAACAAATATAAACAGTAACAAAG
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CTGTGGGAATGGCAATCATAGTATATAGCCCTTACCACCAGCACTGCCTTCTCACATGATTTATTGCAAATTGGTTATA
TTGGAGAGTGGATTCTGCCAAATGGGTAGCCTAATTAGTAGTCAAATTGTTGTTTTCTCTAGTGATTTATTATCAAGCG
TATTAATTTCAGTGTCCAACAGAATGAAATAAATGTCTTGGACTATTTTTTCAATGACATTAAAAAATGAAAACTAACC
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TGCTATGATTGGCGGAGAAACAGTTGCTTGGTAACAAGGGTATACTCCTAGATTACAGCTTGCTATGTATGGAGACTGC
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TCAACTCATGATCCAAGATAGCTGCTAAAGAACTAGATATCATCTCTGATTTCCAGCCAGCAGAAAAGGGAGAAGTACT
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TAGCAGGTGTAAGCTGGAATTCTTTGGCACACAGCCTCTGAACCTTGGCTCATGTTCTCAGATAAATGCCTATAGCAGC
TGAAGCTCTTGAAAAAAATTTAAGAGGAGGAAACAGAGTAGGGATCTTTTTGGGGTCACAGTCTAGGTGTTTTTAGAGC
CACTCAAACTCGAATCAAGACCCTTTGATTACCATAAAAAAATCTAAAATAAGATGTTTTGTTTGGCCCAAACAGTTTTA
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Fig. 6.159

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GAAAAAAGTGAGCCAAATCTGAAAATTAGTTTATTTCATATATATTGAATAGTTAGGCCATTTGGCCACACCTGTCCTG
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CACCTAGGAGCAGCTGGTGTCTCAAGACTGACTGATAGGCTCGAACCTGAGCCATGGTTTACAAAGCTGACACCAGGAT
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TGTTATCTGAATTCATGAAGTGGGCATTACATTGTTTTAAACAAGAAGTATACACAATTGCTTCCCAATCCACTCTCCT
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CCTACTGATGAATGGTGCAAATAGGAAAAAGGAAAATAAAAAGTATTTGCTACCTTGAGAAATACACTCAGTCTCTATT
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TTAACCTCTTTTTCCTAGTGGTCCCCAATTTCATTCTTAGCAAAATAAAATTACAGATTCTGTTTTCAAAGAAAATTTG
TATATGGGATTTCTTTCCATTCCTTATCTAATGTTAGGATACTGAAGTTAAAGTATGCATTTTCTGTTTTTATATATTT
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GAAAATTGTAGCTTTCCTTTCTGCCTATTGCTTTCTGAATGTCAACAGATTGCTTTCTTCAGCTTATGTTAGAATGTCA
CCAGCAGGACCCACATCTGACATGCTCTGGACTGTCAGAGCCACTCAGCACTAGGAGAACTTTCCAGTTGAATTTCTCT
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CCTGGCAAATTCAAAATCGATTACAAACTCATGCATGTTGCATATGTATTTTTAAAGATTTTTACAAACCCAAATAAAA
TAAGTAGAAGAAAACAGTAGTGATAAATTTAAAGTTCTTAACTGAATGAGTAGCTTTGAATTTTACTCAACGGATATAA
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Fig. 6.60

GCTTTGGTTTTTTTTATAAAAAAGAAAACAGGTTTGTTAAGAGACTATGAACATATTTTAGAAATGCTTCAGTGATTTT
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Fig. 6.61

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```

Fig. 6.62

GCAGCCAAATTGTGTTGTGGAGGAATGGGACTCAGGAAGCACGGGCACCCTGAAATAGGTGGATGTGGTCTGTGGAAAA
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Fig. 6.63

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CCTAAACTGCCATATAGCAAATATATAGTGAAATATTATACAGCAATGAAAATGAAACAGTAGTTTTACTGCTATAGGC
AACAATATAGATGAATATAATAAACATAATGTTGACTAAATCAATTAAAAAAGAGTACATACTGAATGACTTGATTCCA
TTTGTATAAAGTTCAAAAATTGGTCAGATAATATTGCCATTTGGGAGATAATAACTGGGATCCTGTCTTTTTCCCTTGG
AGAGCTAGAAGAATGGACTCTTCTCAGAAAGGTGACCAAAACTTCATAAAATCTCCTTTCCCCAGAACATTATTATTTA
CTTACCTAGGTGCTGCCAACCACACCCCATTGTGTCCCTTTAGTCATTGTATGGAAGGAGGTCAAATGCCTTAGTCTCC
CAGTGACTAGAGCTAGATATCCAGCCATATGAACAGTTTGTTAAAACCAGGACTCTGAATTCTAATGCCCAATAAGGAG
GGTTGTACTTTTGTGAAAATAGGCTGGTGACAATTATCTGAGGTCAGAGGAAAGATTCTGTCATCATCATTGTCATGAT
CATAATTCTTATTTTGGTTTCTTTACTCATGTTTTTAAAAATTATATTTGTAGAAGCTATTATGTTAGGTTTAATAGAA
ATGTAGAAAAACATAAAAATCATTTGCTCAGTGTGATGAATTTTCATATACTGAATATATCCATGTAACTTCCACATA
AATGTTTGCGATACTCATCTATATTGTTGCCTGTAACAGTAAAACTACTGTTTCATTTTCATTGCTGTATAATATTTCA
CTATATGTTTGCTATATGGCAGTTTAGGTATGCATTCTTCTGTTGATGGAAACTTGGACTGTTTTCCATTTGGACTATT
ACAAATAATGCTGGTAGAAACATTTTTGTAGATTTGTTTTGGCTCGCACATACACGTTTCTGTTTGGCATATACCAGAA
ATTAAATAGCTGGTACATAGGTAAGCATATGTTCTGTTATTTAATATTGAAGGATGGCAGAATATGCACCCCAGAATAT
GCCATTTTGGTATAAGGATTAGTTTGAACTAAGGGCACTTGAATAACAGCAGATGCAAAAAGGGCAGTCTAATCATTTT
CTTTTTCTTCCCCCTGAAAACAGGAGATAAAAACTCACATGTGAAAGTTGCTCTCCTACTACCAAGAGAAAAGAAACAT
TCTTTGACAAGGAGTCATAGCCAGAAGAATCCTGTACACACAGACCTTGTTAAAATATTCTTATTGATCTTCCTTTAGT
CTTCACACGTAATTTAGTTACTTTTCCACAATTGCCTCTCTTTGTTCAACCAAGTATAACAGCATTTAGGTTTGGCCAC
TTGTTTGGGTCTTCATTTCCTTATGAGGGCTCCTGTGTCATGTAAAAGTTATGTGCTTTTCTCCTATTAATCTATCTTA
TGTCAATTTCATTATCAGGTCTACCTGGGACTCTAAGAGAGCAGAGGTGAAATTTTGCCTCTGCTACAGTAGGTTCTAT
```

Fig. 6.(64)

TAAACGGTTTTCTATGGCATGGGATATGTAAGCAGGGGATAAAGAAATTGTAATAACTCTCAGACTTCTTAGGGAGTTA
GGAAGCTTCTGAAAAAGACCTATCTAAAAGTAGGAGAGAAAGCCTGGGAACTTTTTACTGTGTCTACCTCAGTGAAAAT
ACATAGTTGTTTGTCCTCTATAATTTGATTTAATGTCAATGCAATCTAAACTTGAACGACTTTAAAAATGACCGAAGCC
TGCAGTTCTAAAGATAGAAGCAGAAGTAAGGTATAACTGATGGTAAAACAATATAATATCTGTTAATAGTGAGAAACAA
TCTGTGAGTTCTTTAACCAGTTAAACCTGTGTCAGTGAGCGAATAGCTCTGTGTGGGCAAAACCATTTGCTTAAGAGAA
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AATTGGGTCACTCGACTGGTTGTTCATCCTTACTTAATTCTTAAAATTTTCTGATGGCCTAAGAGTGAAAGTTTCTGAA
TTTAGCTGATCATATTTTAGGAAAAACACAGGACAAAATTTAAGTTGCTGAATTAAAGTGTCTAGTGTTATCCCTGATT
TTTAGGCAAGTCCTTATACTTATCTTTTACCTTTAAAGGAGAGTCAGAGATACCTAGCAGAATGGTTAAATAAAAAGTA
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GGCACTCTGGAAAAGAGTTTGGCAGTTTCTTAAAAAACTAAACATGTAAGTACTACACTATCCAGTAAATGCACTCCTG
GGCATTTATCCCAGAGAAATAAAGACTTCTGTGTTCACAGAAAAACCTGTATGTAAGTGTTTATAGCAGTCTTATTTAT
AATCGCCTAAAACTGGAAGCAACACAGATGTCCTTCAGTGGGTGAATGATTAAACAAACTGTGGTACATCTACACCACG
GAATACTACTCAGCAATAAAAGAAACAAACTACTGATACATACAATGATCTAGATGAATCTTGAGAGAATTATGCTGAA
TGGAAAAAGCCAATCCCCAAGGGTTGCATACTATATGATTCAAATTATATAACATTGTTGAAATGACAAAATTATGAAA
TGGAAAACAGATTAGTGTAGATTAGTAGTTGCTAGGGATTAAGAAGGGAAGGAATAGGAGGAAACTGGGTGTGGCTATT
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AATTATCTCAAAATAAAAAAATAAAAAATAAAGTACAAGAGAGTCAGGGAAAGTTCTATGTAGTTTCTCTTAGATCAGT
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CTGCTTAAATAAGTGGGAGGTTGGTTGAGGTTATTTTCCCCCTATAATTGCTTTTCTTCTGTCCCAGAAGAACCTGTCA
GACCAGTCATTAAGCTATTGTTTAGTTCTCATTTGGCTTAGGTGGTTCAGGGATCAGCTTAGGGAATAATAGAGGCAAT
GTAATGAATCTTAGTGGGAATAAAGTGATTGAGAATTTTGGTTTGTTTTTAAGGCTTAGAATATAATATCCATTCTCTA
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TTTTTCCTCTTCTATGTCATTGTGCACAGGATCTTCCAGCTTTTGGGTTACCATGTCTTTGAGTTTTCTAGTCTGAATA
AGATCTCTTTAGAAATCATATTTTTCTGAGAGCTATAGTTGACTATACTTTGTTTTCTGTTGTTTCATAGGATTATCTC
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AAGAGAGGTAAAGGTGTGAAAAGAGCTGAAAAAGTTTGAAATATGTTAGGGTTTTTGTGTGTGTGTGTGTGTGTGTGTGT
GTGTGTGTGTGTGTGTGTGTGTGTATGTGTGTCCATATATATTCAATGTTTAGCTCCCACTTGTGAGAACATGCAGCAT

Fig. 6.65

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TTGGTTTTTCTGTTCCTACATTTGTTCACTTAGCATAATGGCCTCCAGCTTCAACCATGTTGCTGCAAAGGACATGATCT
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AGGTTGAGTCCATGTCTTTGCTATTGTGAATAGTGCTGTGATGAACATATGCATACATGTGTCTTTATGGTAGAATTAT
TTATATTCCTTTGGGTATAAACCCAATAATGGGGTTGCTGGGTTAAATGGTAGTTCTGACTTAATTTCTCGGAGAAATT
ACCAAACTGCTTTCCACAATGGCTGAACTAAATTACATTCTCACAAGCAGTGTGTAAGCGTTCCCTTTTATCTGCAACC
TTGCCAGCAAATTAAAAAAAAAAACAGTATTTTTTGACTTTTTAATCATAGCCATTCTGACTGGTGTGAGATAGTATCT
TATTGTGGTTTTGATTTGCATTGGCCTAATGATTACTGATGTTGAACATTTTTACATATTTGTTGGTTGTGTGTATGTC
TTCTTCTGAAAAATGTCTGTTCATGTCCTTTGCTCACTTTGTGATGGGGCTGTTTTTTGCTTGTTAATTTGTATAAGTT
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TAGTATATTGATAGTTTCTTTTGCTGTGCAAAGCTCATTAGTTTAATTAGATACCATTTGTCAATGTTTTATTTTTGT
TGCAATTGCTTTTGGCATCTTTGTCATGAAATCTTTGCCAAGACCAAAGTCCAGAATGGTATTTTGTTGGTTATCTTCC
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GGTCCAGCATCAATCTGCATTTGGCTAGTTAGTTATCCTAGCACCATTTATTGAACAGGGAATCCTTTCCCCATTGCTT
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GCCTGCCTCTGTAGACTACACCTCTGGGGGCAGGGCATAGCCAAACAAAAGGCAGCAGAATCCTCTGCAGACTTAAATG
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CCTCTGCTGCTGATACCCAGGGAAACAGGGTCTGGAGTGGACCTCCAGCAAACTCCAACAGACCTGAAGCTGAGGGTCC
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CAAAGGTAGATAAAACCACAAAGATAGGGAAAAAACAGAGCAGAAAAACTGGAAACTAAAAATCAGAGCACCTCTCCTT
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.GAATGAAATGAAGTGAGAAGAGAAGTTTAGAGAAAAAAGAATAAACAGAAACAAACAAAGCCTCCAAGAAGTATGGGAC
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AGCCAAACTAAGCTTCATAAGTGAAGGAGAAATAAAATCCTTTACAGACAAGCAAATGCTGAGAGATTTTGTCACCACC
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AGGATCAAATTCACACATAACAATATTAACCTTAAATGTAAATGGGCTAACTGCTTCAATTAAAAGACACAGACTGGCA
AACTGGATAAAGAGTCAAGACCCATCAGTGTGCTATATTCAGGAAACCCATCTCACGTGCAGAGACACACATAGGCTCA
AAATAAAGGGATGGAGGAAGATCTACCAAGCAAATGGAAAGCAAAAAAAGGCAGGGGTTGCAATCCTAGTCTCTGATAA
AACAGACTTTAAACCAACAAAGATCAAAAGAGACAAAGAAGCCCATTACATAATGGTAAAGGGATCAATTCAACAGGAA
GAGCTAACTATCCTAAATATATATGCACCTAATACAGGAGTACCCAGATTCATAAAGCAAGTCCTTAGAGACCTAGAAA
GAGACTTAGACGCCCATACAATAATGGGAGACTTTAACACCCCACTGTCAACATTAGACAGATCAACGAGACAGAAAGT
TAACAAGGATGTCCAGGAATTGAACTCAGCTCTGCACCAAGCAGACCTAATAGACATCTACAGAACTCTCCACCCCAAA
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TAGCAAATGTAAAAGAACAGAAATTATAACAAACTGTCTTTCAGACCACAGTGCAATCAAATCAGAACTCAGGATTAAG
AGGCAGAAATAAAGATGTTCTTTGAAACCAATGAGAACAAAGACACAACATACCAGAATCTCTGGGATACATTCAATGC
AGTGTGTAGAGGGCAATTTATAGCACTAAATGCCCACAAGAGAAAGCAGGAAAGATCTAAAATGGACACCCTAACATCA
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Fig. 6.6

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CAATTAAAAGAACTAGAGAAGCAAGAGCAAACACATTCAAAAGCTAGCAGAAGGCAAGAAATAACTAAGATCAGAACAG
AACTGAAGGAAATAAAGACACAAAAAACCCTTCAAAAAATCAATGAATCCAAGAGCTGGTTTTTTGAAAAGATCAACAA
AATTGATAGACCACTAGTAAGACTAATAAAGAAGAAAAGAGAGAAGAATCAAATAGATGCAATAAAAAATAATAAAAGG
GATATCACCACTGATTCCACAGAAATACAAACTACCATTAGAGAATACTATAAACACCTCTATGCAAATAAACTAGAAA
ATCTAGAAGAAATGGATCAAGTCCTGGACAAATACACCCTCCCAAGACTAAACCAGGAAGAAGTTGAATCTCTGAATAG
ACCAAAAACAGACTCTGAAATTGAGGCAATAATTAATAGCTTAGCAACCAAAAAAAGTCCAGGACCAGATGGATTCACA
GCTGAATTCTACCAGAGGTACAAGGAGGAGCTGGTACCATTCCTTCTGAAACTGTTCCAATCAATAGAAAAGAGGGAAT
CCTTCCTAACTCATTTTCTGAGGCCAGCATCATTCTGATACCAAAGCCTGGCAGAGACACAACAAAAAAAGAGAATTTT
AGACCAATATCCCTAATGAACATCAATGCAAAATCCTCAATAAAATATTGGCAAACCGAATCCAGCAGCACATCAAAA
GCTTATCCACCATGATCAAGTCTGCTTCATCCCTGGGATGCAAGGCTGGTTCAACANACGCAAATCAGTAAACATAATC
CAGCATATAAACAGAACCAATGACAAAAACCATATGATTATCTCAATAGATGCAGAAAAGGCCTTTGACAAAATTCAAC
AGCCCTTCATGCTAAAAACTCTCAATAAAATTAGGTATTGATGGGACGTATCTCAAAATAATAAGAGCTATCTATGACAA
ACCCACAGCCAATATCATACTGAATGGGCAAAAACTGGAAGCATTCCCTTTGAAAAACTGGCACAAGACAGAGGGATGCC
CTCTCTCACCACTCCTACTCAACATAGTGTTAGAAGTTCTGGCCAGGACAATCAGGCAGGAGAAAGAAATAAAGGGTAT
TCAATTAGGAAAACAGGAAATCAAATTGTCTTTGTTTCCAGATGACATGATTGTATATCTAGAAAACCCCATCGTCTCA
GCCCAAAATCTCCTTAAGCTGATANGCAACTTCAGCAAAGTCTCAGGATACAAAATCAATGTGCAAAAATCACAAGCAT
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GGATACAAACAAATAGAAGAACATTCCATAATCATGGGTAGGAAGAATCAGTATCATGAAAAATGGCCATACTGCCCAAG
GTAATTTATCGATTCAATGCCATCCCCATCAAGCTACCAATGACTTTCTTCAAAGAATTGGAAAAAACTACTTTAAAGT
TCATATGGAACCAAAAAAGAGCCCACATTGCCAAGTCAATCCTAAGCCAACAGAAGAAAGCTGGAAGCATCACGCTACC
TGACTTCAAACTATACTACAAGGCTACAGTCACCAAAAACAGCATGGTACTGGTACCAAAACAGAGATATAGACCAATGG
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GGGGAAAGGATTCCCTATTTAATAAATGGCACTGGGAAAACTGGCTAGCCATATGTAGAAAGCTGAAACTGGATCCCTT
CCTTGCACCTTATACTAAAATTAATTCAAGGTGGATTAAAGACTTAAATGTTAGACCTAAAACCATAAAAACCCCAGAA
GAAAACCTAGGCAATACCATTCAGGACATAGGCATGGACAAGGACTTCATGTCTAAAACACCAAAAGCAATGGCAACAA
AAGCCAAAATTGACAAATGGGATCTAGTTAAACTAAAGAGCTTCTGCACAGCAAAAGAAACTACCATCAGAGTCAACAG
GCAACCTACAGAATGGGAGAAAATTTTTGCCATCTACTCATCTGACAAAGGGCTAATATCCAGAATCTACAATGAACTC
AAACAAATTTACTAGAAAAAAAACAAACAACCCCATCAACAAATAGGCGAAGGATATGAACAGACATTTCTCAAAAGAAG
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ATAGGACCACTTTTTACACTGTTGTTGGGATTGTAAACTGGTTCAACCATTGTGGAAGACAGTGTGGTGATTCCTCAGGG
ATCTAGAACTAGAAATACCTTTTGACCCAGCCTTCCCTTACTGGGTATATACCCAAAGGATTATAAATCATGCTGCTAT
AAGACACATGCACACGTATGTTTATTGAGGCACTATTCACAATAGCAAAGACTTGGAACCAAGCCAAATGTCCAACAA
TGATAGACTGGATTAAGAAAATGTGGCACATATACACCATGGAATTCTATGCAGCCATAAGAAATGATGAGTTCATGTC
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GTTATTTCCTTTCCTCTGCTAGCTTTTGGGTTGGTTTGCTCTTTTTTCTGTTTCCTCTAGGTGTGATGTTATGTTGTTAA
ATAGAGATTCTGGTATCTTGTATCTTTGTTCTCATTCATTTCAAATAATTTCTTGATTTCTGCCTTAATTTCAATGTTT
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Fig. 6.67

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ACCCAAAAGTCATTCAGGAGCAGGTTGCTTATTTTTCATGTAATTGTATGGTTTTGGGTGATTTCCTTAGTACTGATGA
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GGTGGAGGTTATTGGATCATGGAGGTGGTTTCCCCCATGTGGTTCTTGTGGATAGTGAGTTCTCATGAGATCTGATGGTT
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CTTTCTGCCATGATTGTAAGTTTCCTGAGGCCTCCTAGGAGTGCAGAACTGTGAGTCAATGAAACTTTCTTCCTTTATA
AATTACCCAGTCTCAGATATTTCTTCATAGCTGTGTGAGAATAAACTAATACTACTGATTTCTATTTTCATTAAGCTGT
GGTCTGAAAGTGTGGTTGGTATGATTTCCGTTTTTTTGAACTTGCTAAGAATTGTTTATGACACATTGTGTGGTTGAT
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TGCTTGGTATAATTTTCTTCATTCCTTTACTTTGAGCCTATGAGTGTCACTGCATGTGAGATGGGTCTCTTGAAGATAG
CATACAGTTAGGTCTTGTTCTTCATTCAACTTGCCACTCTGCCTCTTTTAATTGGGACATTTAGCCCATTTACACTCA
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GCTTTATAGTATCAGTGATCTATGTACTTAAGTGTGTTTTCATGGTGGTGGGTAATGTTCTTTTCTTTTCATATTTGGT
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TTCTCCTTTGTTTATGAAACTTAATTTAGCCAGATATGAAATTCTTGGTTGGAATTTCTTTTCTTTAAGAATGTTGAAT
ATAGGCCTCTGGCCTCGTCTGGCTTGTAGGGTTTCTGCTGAAAGGCCTGCTGTTAGCCTGATGGAATTTCCTTTGAAGG
TGACTTGCCAGCCGGGCGCGGTGGCTCACGCCTGTAATCCCAGCACTTTGGGAGGCCAAGGCGGGCGGATCACCTGAGG
TTGGGAGTTCGAGACCAGACTGACCAACATGGAGAAACCCCGTCTCTACGAAATACAAAAATACAAAATTAGCCAGGCG
TGGTGGCACATGCCTGTAATCCCGGCCCTGGGGAGGCTGAGGCAGGAGAATTGCTTGAACCTGGGAGGTGAAGACTGTG
GTGAGCTGAGATCGCTTCATTGCATGCCAGCCTGGGCAACAAGAGCAAAACTCCACCTCAAAAAAAAAAAAAAAAAGACT
TATTTTAATGAGTTGGACTCAGATTAGCTAATAAAAATTGCCTAAGTGTGAACCTTCATATTTCCTATGTATGGGGAAT
GTTACGAGGTATGCTGGTGGGAACACAGGAGAGCACATGGAATGATTATCCCCCACTTGCAGAATTTCTTGTTTGGCTA
GAAAGGGAACATACTTGTACAAAAGGCATAAAAAGATGTTTAAAGAGGTGAGCAAAGGCTCAGAGAGTAAGAGAAAGTA
GAGACTACAGAAATATTCCATAAAGGAAGATTCAGGAAGAAGTGTGTCAGAAGTAGAGCAAGAGCGAAGGTTGGATTTG
CTGACTACTACTAATCCATTTCTCAGGAGCATGTGCAGATGCACATGGGAGGTTATAATGTCGGTATCTGTTAGAACCC
TATTCTGTTGGCCAGAAGTCAATCACACAACCCCACTTTTCTGCTAGGGTGGCTGAAAAATACTTTACCTGTGTTCCTG
GAAGGAAAGTGAGGCTTTTTTGAGCATTTAACCAGTCTCTGCTACACACCCACTGGAACTTTGCAGAATAAACTTCAGT
ATTTATTTTTGTTCCCAAACCCTCTACTATCCTGGCCATCATAAAGCAAACATTCTTGATTTTTATTGTTCTTCTCAAA
ATGAGATGCTCACTGTAGAACGTAATACTGTATATGTGATTTTTTAAAATGCTGGTAAAGTGGGATTATTGTTTTCATT
GATTTCATAACTATCAGGTCAAAGAGAGGGCCAAGATTGGGGACTTTGCATTTTTAAAAAAAGAGAATTTGAAGAAGGA
AAAGGGTCTCCAGGATGCCACTTCAGTATGGCAGCTGTGGGAAGCACTAGTGGTAAGGTGCTAGCTATTCTTTCCTACT
TCTCCAAGAGGAAAAAGGTACTCACACTCACAAATCAATTCATTTCCATCAAAACTCCAAAGTGAAGTCAAGGAGAAT
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TTTATTGTGATGTTAACTAGTCCACATGTCTTACATAGTATCTAGGTCTTATGCTTAATAATTTGGCCAGTAGTCCAAA
TTATTACATGATATATAAGCAAATTAGCATTGCTATAAGAAACTAAAATTTATCACCTGCTGCTAATATGGCATTGATG
TTGTGCCATTTCAGCATATCCCTATGCCCCATTCCTGATGTTTTCATCTGTAGATCCAGTGAGGAGAGAAAAAAATGA
GTCGTTTTTTCCTGTTCTTTTCTTTCAACAGTTGTGCTTTCAGTGTTCTCTGTTAAGAAATATTATAACCCTGAGGGAG
TGAATGTTCTTATCAGAGAAGAGAACTGGTTGAAAATGATTTAGTTTCTGAATTCCTAGAAGAAGGGAAAAAGCCTAAG
TGTCCATCTCTCTATCTAGTCTACTAATAGGATTACCTATTTCAAATATATATAATTTTTCTAAATTGAGTTTTTAATG
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CATTATACTTCTCTTTTTCTATAAATGTGGCTGCTTTTGAGAATGGAAGTTGTAGGAAACAGAATTCATTGTAATCTGA
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AGAGGCAGATGCTATGCACATAAATGACATTTATTGATTTGAGTATTAGAGACTTTTTCTTGACAGACTCACCCATCAC
AATATTTAAGTAATATTAACAGTTTCCCACCAAGAACTCAGAGTAAACTAAACAGTGAGTATCCAGGCAGAAAGCCATG
TACTTTTAGAGCTCAATATGTTTATAATATATATGAAGGCAAACATAGAGGCAACATAAATACCTATTCTACTTGGTTG
TAATCTGCCAATTTATATTTTTCGGAGAAGAGTTTATCACTAAATCACTGAGAGGCAGAGTAGGATATCTGGGGGAAAT
GGTGGTGGGCGGGGGGCAAGTTTGTTATCTGGAAAAAGATTGGGAGCCAGGGAGGATTTTCAAATTCGAATCCCTCAG
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TCCTAAAGTGCACTTTGATCAAATTTTGCCTTATGTGGCAACATTTAGAAACACAAGCCTTCGTGGGGTTTCTGGCTCT
ACTTTTTACTGATGGGCTGATAATCTTATAGTAGCATTAGTTTTGCCACCAATTACCAAAGATTTGTTATTATTGGTTT
TTAAAATTGGACAATCCTTAAATGTCTACAGCAACGTTAGAAACAGACCCACAAGTGGGTGGGTCAGACTAGTTTGGAA
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TCAGTTTTGCAAAAGATTATACTATAGAACACATTGAAAAAGAGAAGGTTAGCTGGAAAGTCAAACATAATAGAATTTG
AAAATTACTAGGCTTTAAGTTATGAAAATATTTCCCATATGTTGAGACAGAATTTGTATTCTCTTTTTTCTCAATGGCA
TTTTATGAGACTATATACTGCTGCTCTTCTGTATAAATAGAATAGAAACATCTCCATTTTTAAATTGTCATGGAAATAA
```

Fig. 6.68

GTAATGGCCTGTGAACAAAATGACTTCTTTTTGTTTTGTTTTGTTTTGTTTTGATGGAGTTTTGCTCTTGTTGCCCAGG
CTGGAGAGCAATGGTGCTATCTCGGCTCACTGCAACCATAGCCTCCTGGATTCAAGTGATTCTCCTGCCTCAGCCTTCC
GAGTAGCTTGGATTACAGGTGTCCACCACCATGCCCCACTAATTTTTGTATTTTTAGTAAAGACAGGGTTTCACCATGT
TGGCCAGGCTGGTTGCAAACTATAGGCTGTCACTCTCTATGCTTTAATATTTTGGTTAATTTTAAGGGCATAATTTGTG
GGGACATTGTTCTTATTTGCCATTACTTTTCATAGCAAAAACAGCAATTACTTTTGCACCAACCTAATATTTCCTTTGC
TAATCTAGTGTGACTTCGGAACAGTGATAAATAAAGGTTGGGGAAGAAAGGGCTGACTAGTAAATGCAGACAGAGTTGA
GGCAATTTCACAGTTCCATTGGCGTGTTCTATGGGCCATTTCACTTTCAGATGGTTTGTCTGGCTGAATTTAGAGGCTA
ACCTCTGGGGTTTTGCCTTTCTTCTTGTGACCTGAGAATTTGCTGGCATGTACAAGTCAATTTTCTTAATAGCCTTTCT
TGATTAGTCTATAGCTTCTTCAGTGAGTTGATATGTCACATTTACTTATACTAAAATCTTACACTAGACATATTTTGGT
GTCTTTGGCATCAAACCTCTTTTAGTCATTTGTCTTTCTTGTCTAGAGCTAGTGTCATAATATTTTCATTGTTACAGGT
TTCTAATACACTTTAAGATCTAGAATGACATAACTCCCATTGTATCTTTCTCCTTTTATCTTTTAAGAAAAAGAATTAC
CATGGAAAATAATGCTTCTGTTTCTGTGTCGTTGTTTAAAGTATTTACAACTCTTGTTGTGAATTTTTTTCTTTTAATA
TATTCCTGAACTTTATTTATATATTTAATTAGTTATGTTTGCATTTTTAAAAAGTGAGATAGGTGTATCTTTTCAAAAA
TAATCATTGTCAGGTATGGTATTAGGTTTGTAATAGCTATTAGCTATTACATTACTTTTAATGGCAAAAACCACAAATA
GTTTTGCAACAACGTAATATAATGAATACAATAGGCTTCCATTGTTTATCTATGTTGTGGAATATAGTGTAATTCTGAA
ATAACTTACTTCTTATACAATTTTTGTTTTGCTTTTTTGGTTTAGATCGACTTTTCTAAAAGTCTTTTTCTTTTGTTTC
TGCTATATATTTGTTTTCCTATGTTCCGGTTTCGATTTCTTTCTGGATCAGTTTCATCACTTCATGTTTTTCTTACATTA
TTGCCACTGCATTAGTCTGTCACGTTAATCAGAATAGGATTGTATATGGTATTCTTTTGTATATTTTTAATCTTGTTTC
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ATTGAACTTCTCCAGTTTATTGAGTTGAATATTTTGTTATTTTATTTTATTCAATTTTTAAAATGATGAGAACCCTGAA
TGCCATGTGTTTTTTTCCTTTTGAGTCCATCCTTTACCACACATCATTAGATTTTATATAGATTTTTCAGTTGTTCTTA
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CCTTTGCGTTTTTATTTTTAAAGGTTATCTTATGATATTAATTCCACTGGATTATGTGTACGTCTTTCACATACTTATG
CATTCTTGTGGTTTATATAATTTCTATGTTTTATAGCTTTCATCTGCCTCTATCTAATATATGAACAATTTTTGATCAA
CTTTTTACAAAAGTATTTTCTTGATTTGAAGAATCCAAACTTTTAAATGTAACTGTTAGACCAACTTCATTAATTGAAT
TTACTCATATGCTTTATGGACCTTTTAATTTTCCAGCTACTTTTTCTCTTTAAGAGAGTTTTGTTAGACTTTAGTACTG
TATTTCTGTTAGTTTCTTCTTACAGTTATAACTATGTGTGTGTGTGTATGTGTGTGTATACACACATGTATATATATTT
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TTTTTTTTTTTTTAAGTAAGAAGTTTTATTTCTTCCTGGAATCTTGCAGGATTTTTTTTTCTGTTTACTCAGTAGTTTA
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ATCAGGAAAGTTTTCTTGGGATGTGTATTTGGTTATTGTTTCTGTTTTATTTCTCTGGACTCTCCTCTAGGAATACTGA
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GATTCTTTGATACTATAAGAAGTTATACACGGTATCTGTAATTCTCAACCTATCAGGATTGCTCCTCTGTGCTCTGTTC
ACCTTTTCTCTCTCCATTTCTTTTTTTTTTTTTTTTTTTTTTTGAGACGGAGTCTCGCTCTGTCGCCCAGGCCGGACTG
CGGACTGCAGTGGCGCAATCTCGGCTCACTGCAAGCTCCGCTTCCCGGGTTCACGCCATTCTCCTGCCTCAGCCTCCCG
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CCACCGCGCCCGGCCCTTCTCTCTCCATTTCATCAACCAAAGGCTCTACCTCTGCAGACTAAGCCTTTTGGAACATGTG
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TGCCTCCAAGTAAAATTCCCCAAACCAAGAACTCTGGGTCCCTTTTATCCAATAAAGAAAAGAAGGAGTGTTTGCTTTG
GCAGAACATATACTAAACTTGGAACAACACAGAGAAGATTAGCATGACCCTCACACAAGGATTACATGCAAGTTCCTGA
AGCGTTCCATATTTTCTTACGTAATAGTAAAGGGTTCGATTCAACAAGAAGAGATAACTATCCTAAATATATATGCACC
CCAATACGGGAGCACTCAGATTCATAAAGCAAATTCTTAGAGACTTACAAAGACAGACTCCCACACAATAATAGTAG
AAGATTTAACACCTCACTGACAATATTAGACAGATCATTGAGACAGAAAATTAACAAAGATATTCAGAACCTGAACTC
CATACAGCACTGACTCAAAATTAATCACATAACCGGAAGTAAAACACGCCTCGCAAATGCAAAACAACTGAAATCATAA
ACGCCTGTAATCCCAGCACTTTGGGAGGCCGAGACGGGCGGATCACGAGGTCAGGAGATCGAGACCATCCTGGCTAACA
AGGCTGAGGCAGGAGAATGGCGTGAACCCGGGAGGCGGAGCTTGCAGTGAGCCGAGATCGCGCCACTGCACTCCAGCCT

Fig. 6.69

GGGCGACAGAGGGAAACTCCCGTCTCAAAAAAAAAAAAAAAAAAAGAAATCCAGATATAACCACACAATTACATGGCAAT
TGAACAACCTGCTCCTGAATGACTCTTGGGTAAATAATGAAATTAAGGTAGAAATCAAGAAGTTCTTTGAAACTAATGA
GAAAAAGAGACATCATACCAGAATCTCTGGGCCGTAGCTAAAGCGGTGTGTTAAGAGGGAAATTTAGGCTGGGCACGGTG
GCTCACGCCTGTAATCCCAGTACTTGGGGAGGCCTAGGTGAGCGGATCATTTGAGCTCAGGAGTTTGAGACCAGCCTGG
CCAACATGGTTAAACCCTGTCTCTACTAAAAATACAAAAAGTAGCTGGGTGTGTGTGGCACGCACCTGTAGTCCCAGCTAC
TTGGGAGCCTGAGGCAGGACAATCTCTTGAACCTGAGAGGCGGAGGTTGGAGTGAGCGAGATTGTGCCACTGCACTCCA
TCCCACATCAAAAATCTGGAAAGATTTCAAATTAACAATCTAACATCACAACTAAAGGAACTAGAGAATCAAGAGCAAA
CAAAACCCTAAGCTAGCAGAAGACAAGAATAACCAAGATCAGTGTTGAATTGAAGGGGATAGAGACACAAAAAAATCCT
AAAAGAGAGAAGGATCAAATAAACACAATTAGAAATGATAAGGGGGATATCACCACTGACTCCACAGAAATTCAAATAA
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AGTAACCTACTAACCAAAAAAAGCCCTGGACTAGATGGATTAACAGCTGAATTCTACCAGAGGTACAAAGAAGAGCTGG
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TACCAAAACCTGGCAGAGATATAACAAAAAAAAGAAAACTTCAGGCCAATATCCTTGATGATCATTGATGCAAAAGTCCT
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CTGACCAACATGGAGAAACCCCATGTCTACTAAAAATACAAAAATTAGCCAGGTGTGGTGGCATGCACCTGTAATCCCA
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GGAAAGCATTCCATACTTAATAAATGGTGGTGGGAGAACGGGCTAGCCATATGCAGAAAATTGAAATTCGGCCCCTTCC
TTACACCATATACAAAAATTAACTCAAGATGAATTAAAAACTTAAATGTAAAACCCAAACCATAAAAACCCTAGAAGA
AAATCTAGGCATTACCATTCAGGACATAGGTACAGGCAAAGATTTCATGATGAAAATGCCAAAAGCAATTGCAACAAAA
GCAGAAATTGACAAATGGGATCTAATTAAACTAAAGAGCCCCTGCACAGCAAAAAAAAACTATCATCAGAGTGAACAGA
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TCTCTTGTTGTTATTGCTGACACAAACCAAAGAACAAGAATTTCTTTTATGCCACTGATTTGCAAAGGTCGTATAGAAA
ATGAAGGGTCAGATAATGGATGTTAATCATCCTCATTGATGGATACATTGACAGGAAAACATAATTGAATTGCACATAT
TAGTGGAAGATAAAAAACCATTTCACAAATCGTTCTTAGTAGAGAAAAAAATGATAGGAGAGCTAAAATTGATGTGAAA

Fig. 6.70

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AACACAGAAGGAAGTTAAGGGATTAGAGTGGGAGGGAAATTAGTGAATATCTATTTGTGTGCATGATTGCATTTTTGAA
ATAATTAGTTGCCTAATTTGTGTATCCAAATATAGTAATTGTTTTTTCCATAGTCATGACTTAAATATTCTCTGTGTGTA
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ATGCAAATAAATTTTCTGTACCTGATAAGTTAATTGGAATTTTTTCCTGTAATCTTTTTTTTTTTTATTTTTCTGGTAG
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TTTGCTCAAGTAGAGGGATTATCTGAAAGTACAAAAAGAATTTGATAATTGAGAAATACATTATTTTGAATGCACATTT
GAAGTAGCTAAAAATTCTACGCCACAGCTGAATGCACTGGCAGTTTGTTAATCACATCCATAAACATGAGTCTTCTACA
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ATATTGTGTTAACATTTCCACAATGAAATCTTATTGGCACTTCCTGTATAAGTGGCTGGGAATAGAAATTCTCATTTAT
GAATCAAAACACTTTGCCTTTTTATTTGGAGGTCAACCTACTAACCTTTGGTTATAGGATAATACCAGCAATTTTATTA
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TTTCTCCCAGAAGATTGAGTTAAAAATGTTTTGGTGATGAAATTGAGATGATGTCCAGATTTATAGCATGTTTGGTAAC
CATAACATACTTCTCGATATATTGAATTCCATGTAAATGCTTATTTCTTCAAACCCTTTAAATTTCAGAGTATTTTCTC
CCTTCTGTTATTAATCAGTTCTATTCATAGTGGATCTTAGAAAATTATCCAGTGATTAATATTTCTCCATATTTGTAGC
TTTCATTACTTTATATTTCTGTTTTTGCCTAGGCTCCTTGTTACCAGCCTGTTCTAATGTCACAAAACTTGGCAAACCC
TACCCAAGAAGGGTAAGGTTTAGTATCTATGAATTTTGAAAATACTTGATGAGAAAAGTTGCTTTTAAATTATTTTGAA
ACTCAAACCTTTTTTTATAAATTATACTTTAAGTTCTGGGATACATGGGCAGAACGTGCAGGCTTGTTACATAGGTATA
CATGTGCCATGGTGGTTTGCTGCACCCATCAACCCATCATCTACATTAGGTATTTCTCCTAATGCCATCCCTCCCCTTG
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ATGAGTGAGAACATGTGGTGTTTGGTTTTCTGTTCCTGTATTAGTTTGCTGAGAATGATGGTTTCCAGCTTCATCCATG
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AACTTTTATTACAACTGGAAAGCACTTATGGTACTTATTAGAAAGAAAGTGTGTGTGGATACGTATGTGTGTAAAAATA
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TGTTCCTATTTTATGTAAGCATTGTACCCAGAAGAATGGGTCAGAACCTCTCCTATGGTCTGAATGTTTGTGTCCCTCC
AAAATTCATATGTTGAAGTCCTCACCTCCAAGGTGATTGTATTAGGAGTTGAGGCCTCTAGGGGGGTGTTTAGGTGATCC
TCATGAATAGGACTGGTGCTCTTATAAAAGAGGCCCTAGGGATCTCCGTTAACTCTTCCACCATGTGAGGACACCGTGA
GAAGGTGCCATCTATGATGAAGCTGGCCCTTATTAGACATCAAATTCTCTGGTTCCTTGATCTTGGACTTCCCAGTCTA
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GTCCCCATTCAGGCAGCTGCCTGATGATGTTAGCTGTAATCTCCTGTGTTTGTGCCTGACTTGCTGCCTCAGTTTAGCT
CAGTGGTTCTCAACTGGGGGCAACTTTGACCATAGGGACATTAGGCAACGTCTGGAGACATTGTCACAGCTAGAGGAAG
GGTGCTACTGCCATCTAGTTGGTAGAAGCCAGGGTTGCTGCTAAACATCCTGCAGTGCACGGGACAGCACCTCCTCCCA
TCCCAGCAAGGAGTCATCTGTAGGAATATTTGATAAAAGCCATCTACATAAAAGCCCAACAGCTCACATCATAATTAAA
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TAAATTAGACTGCTCATCAAAAGACATACGTAAATGAGTAAGAGAGCCACTAACAGGACAAAAATATTTGTAAAACATA
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AAAGGTTGAAAACACCAAATACTGGTGAGGAGTTAGAACCACTGAACTCTTACACTTGTTGATAGGAAAATTAAATGTT
GTAACTACTTTGAAAAATGTTTTGCAGATAATAAAAATGTTACTTTCACCTACCCTTTGACCTAGCAATTCCACTCCTAT
TGTTTACCCAGAAGCAAATTTTATGTTACACATAAAAAATTATATCATACATAAATACTTGTTAAAATATTCATAAACC
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Fig. 6.71

CACATATTCATAAACCCACATATTAATCAAAAGAGAATCAATAAACAAATTTTGTCACAGTTATACAATGGAGTATTAC
TCCGCAACAAAAATGAATGAACTACTGATACCTGCAACAAAATGAGTGACTCTCACAGACAAAATGCTGAGTCAAGGAA
TCGAGACAAAAAGGAATACATGTGGTATGATTCCATTTCTAGGAAGTTCTAGAACAAAACTTAGGTTAGAAAAAGGGAA
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Fig. 6.72

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```

Fig. 6.73

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Fig. 6.75

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AACAATTAAATCATACAGAGTGTTTGTTCTTTGACCACAGTGGAATCAAACCAGAAATCAATAAGAGAAAGATAATAGA
AAAATCTGTAAACATTTAGAAACTAAACAACAACAGACTTCTAAAAATCCATGGATTAAAGAGAAACTCTCAAGGGAAA
TAAAAAAAAGTTAAGCTTAACAAAAGTACAAATGCACTATAAAATTTGTGGGACACAACTAAAGCCATGCCAAGAAAAA
AATGTATAGTACCAAATACATACATTAAAAAAAGAGAAAAAGACTCCATCAATAACCTAAACTTTTACCTCAAGAACCTA
GAAAAAGAAGGTAAAATTAAACCCAAAGCACACAGAAGAAAAGAATTGGTTTTGTAGGCTGACTGGGGAAAAATAATT
TATACAACCTTGCTTCTGTTAAAAAAAAAAAAAAACATGAGTTCAAAAAACACATTGTGGCCTTCAAAATGCCAATACCCTAT
TAAATTACCAGAAATTAAAATTAATAAATTGATTGATGGTTACTGTAATGATTAAGATAATGGTTACTTGTGGTGACCT
TTTTTCCAATCATAACCAAACATATCTCTGGAAATGAGTCACTAAGGATGATGCAGCATCTTTCTTTCTCTTTTGATTA
ATCAAAATTATATTAAATATATTTTATCTTATATTAAAGAACAGTTTAGTCATGAATTTAAGCTGGTGTTAGAGGATAG
CTAAGTTGCTTCTCATAGTCATTTCCTTACTTATGTAATGAAGAGCAGAAAAATATTTTTACCAGTGGTTCTCAACCTT
GGCTGGAAATCAGAATCACTTAGGGAGCTTAAAAATATACTGATGCCTGGGTCCCATCCAGCGAGATTCTGAATTGTTC
TATGGCAGAAATTCTCAAAATACAGTATTAGCCTCTCAGCAGCAGCAGCACTTGGAAACTTGTTAGAAATGCAAAT
TCACAAGCCCCATCCCTGATCTAGCAATCTCTGTTTTAACAACTCCTTCAAGTGATTCTGAGGCAGCAGGTCTCAAGCT
TTAATGTGCATGCACATCTCCCAGGGAATCTTAGGAAAATACAGATTTTAATTTGTGGTCTGGATATTACACACTGTCA

Fig. 6.75

CAGATGCTAATACTGCTGGTAGCAAAAATATAATT_GAATTACAAGGGTCCATAGGACATCTGGATATTTGCATTTTAA
AAAATTTTCCAAGTGGGAATGCAGCCAAGGTGAAAACAACTGGTCTAGATAGCTTTATGGTACACTGCCAATAGCCCAA
GCAATCTGAATGATCTCTGCTTGGTTTTCTGTACCTGAGGTTGTAGAGTCACTGAAGAGCACATACTTCTTGTTCTCTT
TAAAGTGATAATCCGGCTGGACATGGTGGCTCATGCCTGTGATCCCAGCACTTTGGGAGGCTGAGGCGGGTGGATCACT
TGAGGTCAGGAGTTCATGACCAGCCTGGCCAACATGGTGAAAACCTGTCTCTACTGAAAATACAAAAATTAGCTGGGCG
AGTGAGTCGAGATTGCACCACTGCACTCTGCACTCCAGCCTGGGTAGCAGAGCAAGACTCCGTCTCAAAACAACAACAA
CAACAACAACAACAACAACGACAAGAACCAAACAAATAACAACAAAATAAAACCTAAAGTGATGATCACTGATTTTAAG
TGGCTCCTTAGTTGCCTAAGAATCCCAGTTGTGATGGTTTTATCCCTTATTGTCTAGAACTAATGTTGAACACCCTGCT
TTTTAACTTCATCTTGTTTTTCTCTACCCCCATCATCAATATTTGCCTGACTCACCATCTTTCAAGGTTTACCTCTTCA
TACTCAGCTAAAAATTAGCTGATGAGATGCACAAATAATTCCAGTGTATCAGGAGAGGATCAGCTTCCTCCCTTTTAAG
GTAACATGGATCCTTTCACCATCTCCTGATGCCCTGAGACCAATGTCTTGAGAGCATCACATCTTGTTTCATGCATTCT
GACTGGGCATGCTCATCTATTCCAGTGTTCCCAAGAAACACTCATGTATTGATCACTCCGTGGCCAGGCCAGATCTCTC
ATCTTCCCTACCATGTCTGCGTCTCCTCCTGCATTCTCTGTTTCAGTGATAAGTAACTTGGGGGGGCATTTTTGATTCTC
CTTCCCTCTTTTCCACCCCAAATGGAACTTGTTGTTAGGATCTGTGGATTCTATCACTCTTGATTCCACCTACTTTTCA
CCATTTTGCCTCTACTTCTCTGGTGCAGGTCTTCACCATCTCTTGCTTAATTGCTGCTACAGTGTCCTAACTAATTTCC
CTGCCTTTAGTCTGTCTGTCCATCTCTAATCCACTTTCCATATGGCAGTGAGAATGTTTATTCTTGCCACTTCTTAGAT
CTAAAACTCTTTAGTAGCTCTCCATTGCTCTCAGAATTGAGAGTCCTTATTATGACTTAAAAATTATTAATTATCTGGT
GCTTGCTCACCTTCCAAGGATCTTCTTTTCACCTTGAACTGGATGCATTTTCCGTATGTACCATTTTCTCTGTAGGTTC
TAAGCCTTTGAAGAGGTGATGCCTCTGCTTGGAGTGTCTTTCCATTCTCCCATTTGCCTAGTTCTTATTCTTTTTTTAT
TTGCTAAGACACTACCTCTGAATAGATGACTTCCTTGACCCTGCCTCTCTCAAGGGTGAGTTACATGCCCTCTTCTTAT
CCTTAGTGTTAAAATTACCACACTGACCTTCAGTTCCCTCTTTACTTGTTTATCTTCCTCAGTAGCTCCTTAAGAAGGG
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GAACTAGATGAGGATTTTTGTCTAACTGATACAATTCCACGTTCTGAGTCCTTGCCTTCTCTCTCTTTCAGTTTCTGGG
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TGGCAGTCTTCTCTGTTGAGTGATAAACCAATGCCATTCTAAAAGATCTGCAGACCTGTGTGATCTCTTCTTCGAGGGA
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TGATCCCTTTATAGACCACATTCTAGCACAAATTAATGTCACTGTTTCTTCTGTAGGAGGAGCCTGGCACACTGAATAC
CAGTGGAGTCTTGCTCTCTAGGCTTATCTTGGGAGCTAGTATGCTTATTCTTGTTTCTGTCCCTTTCCTTATGCACTGC
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ATGTTTTGTTATGATAAAAAGAAGCAGTTCGATCATGACATTTGCAGTTATCTTTCTTGCAGTGACCTTCTAGAAATGA

Fig. 6.76

CTCTTCACCTTCCTATCTCTTTCCCTGTCTTGGTTCATGGTATACTCATTCTTTCAATTGTGAAAGTAGAAAGTCAGCT
GACTCTTTGATTCTTCTGTCTCCCTCTGAACCACTGACTTATGGCAATTTCCCTTTAGCATGTGTTCCGTATCTGGACC
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GACTCGGGGCTCAGCTTGGCAGGAGGTGTGATGTCTCAGAGCAAGTACAGCATTTTTTGAAGGAGCAAGGTGTTAATGG
CAGGTGACTCTGGCCCCTTTTATGTGCTTGAGCTGTTTTGCCAGGTACAGAGTGGGAGTGGAACAGAAAAAGGGTTTTT
TCTAGTCTAGGGTCCCACCAGCTAAGGCCTCCTGATTGAGCCATCTGTAATCCCAGGGTGGACCTTGAGGCAGCCTATA
AATCATTCTCTGCTCTGCCACTGTGTCAGGTTAAGGCTTGGAATTAAGGTTCCAAGAGAGGTAAAAGGTAAAACCAAGA
GCTCATCAGACAGCCTGACCACAAATTTCCCTCTCCATTGTGCTCCTGTGGGTGAGGTCTCCTAGCCAAATGACTTTCC

Fig. 6.77

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GGGAACTCCTCCCTCACCAATGGGGTAAAGAAGACACTGCATTCAGCTTTTCTAGCCTCCTAGCATGTATAACTATATT
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CTTTCTGAGTAAAGATTCCACTAAATAGTTTCTCCTCTCACTTGCAAAATGTTGTCACTCCATAATCAGGGATGGAATG
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CCAGTACATGTGAAAGGACTTTAGTGCCTCCCTCTTTTGTTCTCTGGGCTATCTTCTCCAATTCTGATTTGTTGATGTA
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CCCACTTCTAAAGCTTTGTTTTCATATATTTGCCCATTCCTCCAATCTAAGTCTCACCATTTAAAAAGTTAAGTTCACA
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TCTGTGTATTGAAGCAACAGAAACCATGAAGCATTTTATTGCAAGAGCAATCAGGCAACTTCCCAGATGTTGCTAAACT
CTGCTTTAACTGTTGTGTAGAGCTAGTTTTAACTGCATCTCCCACTGCTACTGGAGACAATGATTCTAGAGATGTTACT
AATTGTCACTTGTTCTTTGCCTAGATTTCTTCTCTGCTCCTTGTGGCGTCACCCAGCCGCTAGACGGGCAATGGATTAG
AATTTACATGGAGGCTGTGGTCACGGTTGTCTCAGGTGTTAAGGAACAAAGAAAAAGAGGTGCAGAAAGAATAAATCAA
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TGGGGGATGTTCAAGGAAATAACAAACTGCAGAAACATCATTTTTATCTGTTTCATGCACACTTGTGGCCCAGGAGATT
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CAGCCCACCTCTTTCTTTCATTATGAGCAAAAGTAATCAAAGACCATCAGTGCTAATTAATTCAGATAAGCCAAAAACA
TACATATTCTTATTTTAAATGAAAGAAATGTTAGTGTTGTGATTGTTTTTATATTATTTTAATTAAAAAGTTAAATGGAA
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GATGTTAATCTATCCTGTTTTTAAATGAAACTTCTAGTACAGAAGACACATGGTCCCTTAGGAACTCATTGTGAAATTA
GAGGGTTTTCAGAGAAGGACTCTGTTGTTTATTTTGATTGTTCATCTTCTCCATAGGGATGATCAACTTTATATAAACA
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Fig. 6.7$ (crossed-out)8
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GCGCTGCATCCATGTAAAGTCATTTTACATGGCACTTGAGCAGATTGACAGATATTTAGAGAACATGACCGCCTTAGGG
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TTACAATAAAATTTTTAAAAATCATAGAAGTCTTAAGTAATATAGTTTTCACTAACAGAATTTTTAATTGTTTCTGCATG
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AGCTGGAGTAGACATTTTTAAACAGGAATATTTTTATATTTTTCCTTTTGAATATTTTCAAAGCATCAAACACTTATTG
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ACCAGCAAAGAACATTTTTTTTACTTTTTAAAAATATTGACTGATTTTAATGGGAAAATGCTTTATTATTTTGTTTAATT
TCTGAGGGTATTTTTTGGTCTTATTACTGAGGTCAAATCTTTTCTCATACTTGTCATTTTTGTTCTTTTGTGTTTAATT
CATTTATGTCTTTCTATTTATTTTTCTTATTGAATTGTATGAATGCTTCATTCCATAAAACATGCTATTTCCTGCAGGG
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TATTATTAGATAGGTCCAGAAATTGCTGGTACGTTGGGGCTGTGGGCACATTTGGTTTATGTTGCTTTCCTAGATGTTA
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TCTGTCAGCATCTACTGGCAGAAGTGCTTTCTCTAGCATGACTTCACCTCTCAAATCCCTGGATGTAACAAAAGTCAAT
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TCTTTCATTCTTTCTTTCTTTCTTTTTCTTTCTTTGACTCTATAATAACTTCTTATTACATGCAGAAGTAAATCT
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CAAATTGTATATATTTAAAGTGTACAACTTGATGTTTTATTCTATGTATTTATTATGAAATGATTACATTCCAGCTAAT
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AATACTGAAATTGTGAGCTCTTTCTTTGACAACAAATGTGTCCAGGGAGATGATGAGCTTGTTAAACATCAAGGTTATA

Fig. 6.79

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ATTCAAATGACACCATCAAGTTGAAGAAATGTAAAAAGATATACAGACACAATGTTTGAGGAATCATTTGCCAGTGTTA
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CTCACTATGTTTCCCAGGCTGGTCTCAAACTCCTGGAGCTAAGTGATCTTCCCACCTTGGATTAAAGGTGTGAGCCACT
GCACCTGGCCATACTTACTTTTGAGTAGTTGGTGATAATGATGATGTTGATAGAGAGTAATTAAGTATGAATGAGTTAT
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CCTTTGACAAACATCAGTGTCTTCACTTTGACAATGGTACGTTTTCTTTTGCTTATCATCCCTGGTTAAATCTTTTTCT
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GAAAAAGTGTAAATATGACACTTCTGAGCCGGGGAAGCAAAGTGTAAACTGAGGCAGCAGTATTCTTCTTTATGAAGTA
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TACAATGACCTTATGATGTATATACCATTAATATTCTCATGTCATAGATTAGCAACCTGAGGCACAGAAAGGGTAAATC
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GATGGACAGTGGGAAGGGGGAGAGGATCAGGAAAATAACCAGCAGGTACCAGGCTTAATGCCTGGTGATGAAATAATCT
GTACACCAAACCCCCATGACATGAGTCTACTTATATAACAAACCTGCACATGTACCCCTAAACTTAAAAGTTAAGAAAA
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Fig. 6.80

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TTTCTTTCTTCAATAAAAAATTAATAGCTAATTTTTATCTTTAACAATTTTGACTCCTGTAATAACACTACTTAAAACG
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TACAGTTAACTATTTTTTTAATAAGTAGAAGGACCACACTCTAAAGTAATGATAAAAAGTATAGTAAACACATAAATTA
GTAACCATTATTTGTTATTGTTATCATGTATTATGTAGTACACATAATTGTGTGTGCTCTAATTTATATGAATGACAGC
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CAAGGGTGATGAATACATTTTATTTCTGTTTTCTCATCCCATTTTCTATTGCAAATGAATTTTTTTTTCAGAAGTAGGTA
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TGAAAAAAAATCATCAGAACAGAAGAGACAAGGAAGCTGTGTCTTAGAAAAGAATGTCTCCTTCAAAAACACATCTCT
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GCTGGAGAGAAAATTAGTAGACTGTAATATAATTTGAAGAAATCAACCAGAAGGCACTGAAAAGAAAATGATAGAAGAT
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Fig. 6.81

ATGAAAAAGAAGTTAAGAGACATGTTGGAGAGCATAAGATTCAATATATGTTTCATAGGCCTTCTAGAAGGAGAGCACA
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CCACATGGATTTGAAATTAAATCAACACACGTCTAAACAACTCATGAGTCAAGCAAAAAAATAATGAAAATTAGAAAAT
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AATTTTCTTATACTTGGATATTGATACTCTTTTCTAGGTTTGGGAAGTTCTTTGTTATGATCCCTTTGAATAAACTTTC

<center>Fig. 6.82</center>

TACCTCCTCTTCATGGCCAATAACTCTAAGATCTGCCCTTTTGAGGCTATTTTCTAGATCTTGTAGGCATGCTTTATTC
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TGAGACCAGCCTGGCCAACATGGTGAAACCACGTCTCTACTAAAAATACAAAAATTAGCCAAGTGTGGTGGCGTGTGCC
TGTAATCCCAGCTACTTGGGAGGCTGAGGCAGGAGAATTGCTTGAACCCTGGAGGCAGAGGTGGCAGTGAGCTGAGATC
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GAGACTCTGATGTATTCTTCAGTATATCAATTGCATTTTTCAACTCCAGAATTTCTATTTGATTCTTGTTAATTATTTC
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ACAGTTATTTTGAATTCTCTGTCTGAAAGGTCACATATATCTGTTTCCCAGGTTTGGTTTCTCATGCCTTAATTAGATT
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ATTATGATCTAAGCCATATCTGCATCGGGGGTACCACAAACCTAGTAATGCTGTGGTTCTTGCAGAATCATAGAGGTAC
TGCCTTGGTATTCTTGCATAAGATCCAGAAGAATCATCTGGATTACCAGGGAGAGATTATTGTTCTCTTCCGTTACTTT
CTCCCAAACACAGAATCTCTCTCTATGTTGAGCTAACCACCTAGGGCTAGGGGTGGGGGTGACAGAAGCGTGCCTGTTGC
ACTGAGACTGCGTTGGGTCAGACTTGAAGCCAGCATAGCACTGGGTCTTGTCCAAGGCCTGCTGTAACCACTATCTGGC
AAGGCCCTAGATACCTGGCCGTACTTAAGGTACTGTTCACTTAAGGCCCAAGGACTCTTCAGTCAGCTTGTGGTGAATG
CAGCCAAGTCTGAGACTCATCCCTTCTGGCTCAGGGCAGGTCCAGAAATGCCATTCAAGAGTCAAGCCTTGGAACTGGG
GACCCCAAGAGTCTGCTTGTTGCCCTAACCCACCATAGCTGAGCTGGTACCTGATTTTTGGTTCTTCTGAAGGTGTGAT
TTTTGCCTAGATAGTTGTTACATTTGGTGTTCCTGTGGGGTGGGGGGACAATCAGTGGGGCCTTCTATTCCACCATCTT
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GAACAAGAGAAGAATGACTTCGGGTTCTGCTTATGTTTAATTGGTTGTTCTAGCCAGCATTGTAGGTAAAAGAAAATAAA
TAAAAGGTAGATAGAAAAGAGAGAAACCTAATTGTTATTAGTAAAGAGACTTCATTGAAAAACTATTAGAAAATAATAC
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ACCAATTGTAGGATTTTTCTGTGAAGGGAGGTAGGGAAGGAGGTAAATGGGCTCAGAGAGGGAAGGATACACTGGGCTT
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TAAACTATTCACCTATTGTGGGACATTTTAGTCTTTTTCAATTTTTGGCTACTACAAACAACTGCATATAAGTTTTGCTG
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TTTAGCAGTTCAAAGAGGTGTGTAGTGATGTCTTATCATAGTCTTAATTTGCATTTCTCTGATGGATAATGTACTAGTT

Fig. 6.83

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TTCTATTGATGCTGTAACAAATAACTGTAAACTTAGTGGCTTAAAACAACATAAACTTACGATTTTACAGTTCTGTAGG
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CATTTGTGGGTTCAGTCCTCATATCATATCATTCTTACCTCTTCTGCCTGGGTCTTCCACTTTTCAGAACCCCTGTGAT
TAAAGTAGCCCCAACCAAATAATCTGAAATAATGGCTGTATTTTAAGGTCAGCTAATTGGCAATCTTAATTCCATCTGC
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CCTATCATAGGTAGATATGTTGAACATCTTTTCATGTGCTTATTTGCCATCCATATGTCTTCTTCAGTAAAATGTCTAT
GTCTTTTTGCCATTTTCAAATTGGATTGTTTGTTTTATTAGTATTGAGTTTTGAGGGTTTAAATTTTTTTGTTGTAGA
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GATGTTAAAAATAAATTGGCTGTATTTGTGTGGGGATGAATTGTTTCATTCCTTAATGAAAAGAGGAAATACAATAAAC
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TGAAATAACCTGCTTTACTCAATTTAGTACACAAGTTATTGAGAAACATCTATAGCCTTTTTCATTGCAGTGTGGGACA
GTAGTTCAGTGCATAGGATCTGAGGCGCTGTCACATCTTTCTTGATGTCTCAAAAACATACCATTGATTTTGCTTCCCA
AGTCTAGATTTACCAACATGTGAGAGACTGATGCTGCAGTGCCAGGAGAATTATTACTGATCATAATCCAAGAAGAACT
GACCAGAGAATTAAAGCATTTAAAGTTGCAAGAAGTAGGTTTATTCCTTGACTCAAAAAGGCTTAGTATAACCTACTTG
CACCCCTGTAAGCATCTAAATTATTGTCTTAAGAGAACAGAACCATAATGATAACCACAGTATTGCTTAATATCTGCTG
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Fig. 6.84

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CTGTTTTAAAGCATTTCCCACAGTATTTCATTGTGTCTTAATGTTACTAAACTTTATATTGCAATTGCTTAGAACTGCT
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ATCCTATCATTGAAAATTCTAGGGTTAGATCTGGATCCCAGGCTAGAAAAATATCATTAAGATCATTTTTTTTTCCATT
GCACAGCTCAGTTTTTCTCTTTATTGGCTTCATTCAAGTAGACTTTACTCCTACAACCCTGTCACCAACATAAATAATT
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ATGTCAACCTCAGTTAAAATCAGTAGTCCACAGTTGTTTTTGCCAAAACAAGAATGAATTACATTGGTGGTGAAAGTCC
CTCTTCTAGTGGCAGAGAGAGAATCTAGAATAATTCACCCACTCGGGATATGGGTGGTTTGCAAGGAATGAAAAATCCA
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GAGGAGGCTAAACTGCCATTAGAATGAAGAAGAGTTATGTAAGACTATTTTTTATCCAAACTGATTGGGCTGAATTCCA
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TTCCTCCTTTCCTCTCCTCCTCCCTTCCCTCCCTTCTACCCTTCCTCTCCTCCTCCCTTCCCTCCCTTCTACCCTTATT
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Fig. 6.85

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TCCCTCTTTCCTATCACACAATCATTTGTTCAATATATACTTCTGGAATGCTGCTGTAGCCAGGGACTCATCCGGGTGC
TAAGGATCTATCAATGAACAAAACAAAGTCCCTGACCTCATGGAACTTACATTCTGGGAATCGTATGAAATATTTCTCA
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CAAATGCCAATCATATATTAGGCATATCCACCTATGTTATGTTATTTAATAACTAATGGCTTACTGTGATTCTGACTGT
CCCATTTAAATTCTGATTGTAGAAACATATTTTGTTTTTGTGACTGATTGGGGAAACTTTAAATATTCAAGAAGTTAGG
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CCTCTGTCAAAAAATAGGTGGTTTGTTCAAAGACTGATTGGAAATTATTTTAATATAGACTGAAGATGACAGACTCATC
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GCAATTCAGTTTTATCAGTGATCCCAGATTTAATCATCTCACAGTCTCTCAGATGATTGAACAAATGAACCTAATAGTG
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TGCACCCCACTGCAAAACCGAGATCCCTGAGGAGGCAGCGATAATAGTCTGAAATGAATTTGAGGCTCCAAGAGCAGAA
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GACTGCCTGTAGGGGAGAGGGAGGAGCTCAGTTAAGTCCTGTCTTCCATCCATCTCACCAAGATGACTGGCATGTGAAT
GAAGCCATCTTGGATTTCCAAACCAGTTCAGCCACAGTGAAACACCACCAAGTGACCCAAGGGAAGCAGAATCACACAA
CCCAGTTCTTAAAAATATGACCCGTGGGGCATGAACATTATAAAAATAATTATTGTTGTGGCCATTAAGTTTTGGATGG
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AACTTCAACCTCAAGAAGTCTTTTGGTCAGAGAAGAACTCACAAACTGACTCCCAAGGGAACACTTGGGAACAATGCTA
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AACTCATTCATTTGCACATCAGCAGTGGGCATTAGCATGGACAAATGTTGACAGAGAAGCAAACAATACTGCCATGATT
GGTATCATTGGAGTTGTACAACACTGCAGCCTTGCTGTAGCCACCCACTTACTCTCTAAGTCTTAGCACTACCCAGGAAA
AAGGGGGGATGGTGAGGAACCTTCATTTCACTGGGTAGCCACCCACTTACTCTCTAAGTCTTAGCACTACCCAGGAAA
ATCATCTTGAACTAAGTTGGTTATTTGCCAGTAGGAATATCCTCATATGGCAGAAGAAACTGTGATTTGTCTGAAGACT
ACAGTTATTTTTTAGCACAATTGAGTTTGCAGCTTGTGAGATCAATATCTACCATAGTAATTAAATATGTGAAGTACAG
CCGGTAAGCCAATTAGAAAATCCCCATTATCTGATTAGCTCATCTCTTCCTGTGTCCATTTATTCTTTTACAGTGGGAA
TGGTTTTTCGGTTGTGATCAGAACCATTGTGTCTGCTCCAGTATGTACTCATGCTAGGAGGATAAATTAGTCATCAGGC
CATTCAGTCATCAGAAGTTCTGAACAAAGGTAATAGGAGAAGCTCTGTCTTGTGGGTCATCATGAGCCTGATGAACCAC
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TTCCATACGTGCTAAATAAATTATAGCAGATTCTGGCTTCCTGGTTACCATTGATTTCCTTCTAATGTGACCATTCCAC
AAGTAATGACAAACTTAATCGATAGGTATGTTTATACTGTTAATTAGGTATACTCAAAGCTTGTTTGGGACTTTGTAAT
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GAGTATCTCACTGTTGCCTCCATGTACTAAGGGTCACTCTATTCCCACCCTGTCCTGCAAGACATTCAGTGCAGTGCCA
ACCACACTTAACAAGAAGGATCCCAGTAAATAAGAAAGACGGAAGCATTTACATATTTTTGCTTAGGAAATGAAAAGAAG
GAAAACTTTTAACATCTCATTATAAATATCTACCACCCAGTAATTGAGTTATAGCTTATATTGTGCAATTTACAATTCA
TATCTTTAAAAGAGCTCACAGTTACCTCTATTTGTAAGTTTTGACAGCAAAAACCTGTTCATTTCTTCATCAAACATTC
ATTAATCCCTAATTAGTACCAAGTTCCATGGTTAATGCTTATTAACAGTCCTGTGAAAGGAATATTAAAAATATATAAA
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Fig. 6.86

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CCACAAGGCCCATAATAAACATATGAAATGTTTACTTTTATTAGTCATCAGAGAGATATAAATTTAAACCTCATTGAGA
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CCCTATGATCTATAAATTCTATTCCTAAATTGTATTAGTCATCTGGATAACAAATTAGCACAATGTTAATGGCTTATAA
CAAGAATAAACACGTATCATTTCAATTTCTGGGGAATCAGTAATTCAGATGAGGGTTTAGGTGGTTTGGACTCAGAGTC
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AAATTAATACAGAAGGGAAAATATTAAAAGGAGAAACTAAGGTACATATATTTTTCAAATAAGTAAACACATGGTAATT
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CATTAGCTTCATTTTTACTGTGATGTAATACTCTATGGAGATTTCATTTTACTCTTGATAGATCTTTATTTTTCATTTTT
CAGTTATGAATGTCTGTTGTGATTTTTGATGAATACATCTACTCATTCATTTTTTAGTTGACATAGCTAGAGTTAACAT
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Fig. 6.82

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GGCTGGTTTGTAGGGTAGACATAATTCAGCTTTACTAAATGCTGCCAAAGGATTTTACAAAGTGATAGTACTAACTTCT
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AATGATTCTGGTGGATGTGTGGGAGCCCTAGTTCCTCTGCATTTGTGCCCATATGTGATATTGTCAGTGTTTTTCATTTT
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CACTATTTTCTGTGATTGTTCATGCTTTTTGCGTTGTGTTTTACTATGTGATTTGAAAATCCAGTCACCACAAAATGCA
TTCAAATTGACAGATGCTAGGTTTGTTTCTTTCCCCATTAAAAAAATTGTAATATTCACATACATAAAATTTACCATCT
TGGCCATTTTTAAGTATAAAGTTTAGTGTTAATAAAAACATTCATAATGTTGTAAAACCATAATAATCATTACCCATCTCCA
GAACTCTTTTTATTTTATAAAACCAAAACTCTGTATTCATTAAACAACAACTCCCCATTCCTCCTCCTTGTCCCCATTGTCT
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GCCAATGACAGAGGCATAAAGGAAGACTGGCTTGTCATTGTGTAAAGGTGTAGCAAAACGCAGCAGCCAAGGCTACTC
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GTGAACCTTTGGAGTTCAAGAACGTATAGTATGATGGAAATCCTGGGAACAGGAGGCTGCCACCAACATGGCTGCTACA
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Fig. 6.88

GCCAGCAACAATAACCAAAGCAATAGGAAAATAGCCCCTGCTTCGCTTCTGTCTGCCAGCTGTCTCACTAGAGCACATG
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AACAAAAATTCTGTGTACCCCTAGTATTAGGCAAGACTTTGTTACAGAAAATCAACTTAAAGTCATTTATGCAAAAAAT
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CAAAGATCTCAGAAATCTGGTTTTAATTTATCAGCTTTGAGTTGCTTTTTTCCTTCACCTTTTTCATGCTTGTCACTGGCT
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ACTGATCATTGTTGCTGCTTATAAGAATACAATGGGAGACAGGATTCTGAGGCTGGGTGCTGGGGGAAGCCATTTAAAA
TCTGGCAGCACTGGCGGGAAAGCATACCTGGTGCTGCCAGGTTGGAGCTTGGTGCCCTTTTCTGCCTTTTTCATGACAC

Fig. 6.89

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CTCTAAGGGGGAGCCATTGTCCTGCTATTCTCTTTGGCTTTCTTTTTAAGGGGCTGTGCACCCATAGATTTGCTGTAGC
CAAAACTTACAAGACCGTAACTACACAGTTCAACCCTACATACACCCTGTGATATCTCTGAACTCTTCTCTACTCCTCT
CTTCCCTGCTCACTTGGCCCCAGCTGTGCTTCAGACACACCAAGGTACTTCCAGACACTTACAGGCTTCACTCCTGCCC
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TGCTAATACACTCTATTTTTGTCTTTGCACATATCACCATCTCACAGAATGTATTTTACTCACTAATCACTTACACATT
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CCCATTTAAGTTTGGGTCAAAATTCAGTATACTGACTTATTAATGAAGTCATTTGGAATGAGTAAGAGTCCTAGGTCTA
TGTCGATCACACTGAAATAAGAGCAGACCATTGATTGATTCAATACTAAATGTGATTGTTCTATAACTTTCTGAACACT
TCATGTAACAGCTCTGTCTAAATCTCAGAATCTCATTTATACCTCATCACTGAATATTTTTAGAATATTGAAGGAGTTT
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ATACATTATTCCTATTTGAACTGAAATTCTGAACATCTTTTTAATTACTCAAGGTTTCACAAATTATGAGAGAAGGTTA
TGATCTTTTCAAGTTATGCTGTAAAATAAGGACTTCTGTTAGAAAACAACAAATAAATGATTAGACTACCCTCCATGTT
ATTCATCCTCTGTTAGCTGAATATTCGTCTTAAAGTAGGGCCCTGCCATGCCATTCTCCTGCTAAGAGTCTGCATTGGT
TTCCTACTGGATCAAATCCAAGCTAAGGCACTCAAGTGCCAAGCAAGGCACTCAAAGCCTCATATAGTCTCATTTCCCC
TTAACCATCTTAGGTTATTTACCTCAGCTGCCTGTAACAAATCCTCTGCTCTAGTCAAGAAGTTTCTCTTCTTCTTTTC
TTCCTTCCTCCTTATTAGCCACCCCTCCTGCGTTAGGCACCCGTTTACTAGTCTGCCCTGCTTCAGGGACTTCTTATTT
CTTCTCTGATGTTCCTTTCAGGCATGAATGTCATTCTTTTTTTCCTGCTTATCCATACACCTTATGGACCTATCTGTAA
GATCCACCTGTAAATTACACACTTTGTTTTCATGTCCCTAGCCTACAACAAATAATCTCTCTCATTTTTCTTTTCTTAA
TTACTTTTGAACTTATCAGGTGTACCACACAATTTATTAATCCTTATTTAATTACAGACTGTAGCATTCTATAATTGTT
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CACCATATTGGTCAGGTTGGTCTCAAACTCATGACCTCAGGTGATCCACCCGCCTTGGCCTCCCAAAGTGCTGAGATTA
CAGGCGTGAGCCACCATGCCCCACCCTATTGCCTTTTCAGGAAAGTTTTTGGAGTGTTCTGAAGGTTGGGGAGGATCCA
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GCTAGCTAGATAGATAGATAGACAGACAGACAGATAAAAGACACCACAGGCTATAGTGAGAGGTGAGACAGCACCTGGC
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```

Fig. 6.90

GGTGGCTAGGGCATAAATATTACATCCTATCTCCAGTTAAGGCATGGATACCTGCACCATCTTTCATATGAGAAGCATC
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AAGTGATGGGTTCATTTTGAGGCTCATTGAATGAGTCCTTGAAGAAGGAAGCATCAAAATGAAAAGGATTTCAAATCTC

Fig. 6.91

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TGAGTATAAGGTTTGAAATGTTCATTTGACCTCACTGATGGCTGTGGAAAAGCCTAGGCTTGGACACAGAAACTGAAGG
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TATCTATCTATCTAACTGTCTATCTATCTATCTATCTATCTATCTATCTATCTATCTATCTATCTATCTATCTATCTGC
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TGAGAAACCGTATGGCGTATGCAAAGAATTACAAGTGGTTCCCTATAGCAAGTGAGTGAAACAATGCTGTGTGGGAGGT
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Fig. 6.92

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CAAAACAATGTGATAGCAGAGAGTTCTAGTCAGATAATTTGTGAAAAGTTTGTTAGATTTTGTGATATGGAGGTCTAGG
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GAGGTAAAGAAAGATAACAGACATTTTTTTTCAAATAGCTTGATTTTGAAGAGAAAGAGAATGATATCTTGAGGAGGGA
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CGGAAACTAGCACAGGCAGGGAGGCCTCAACACCTAATTTATTTCCTAGGTGAAGTGGAACCTGAAGATTTTCCTTTTA
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ATCCTTGTATGTTTCTTTTCTTTTAACTTTTTGAGTATTTAATTCTTGTGTTGCCTCCTCCTAATTTCCTTCCTGCAGA
TACTTTGGTGAGTAAACCTTGATAAATTAAGCCTGTTTTTATTATATTCCTTTGGAGTTGTCTTTGATTGTGATCAAGC
TTCTCTTTTTATACATGCCACATGTATTCATTCTTCATTTAGATGAGAATCTAATCAAGGGAATAAACTGCCAAGTTTG
GTTTCATTTATGCAACCCTAGAAAAATATATCTTTTGATGAGTTGGAGACAACAGTAAGTTAGAGACAGAGCTAATCCA
TTACACCTTGATCACTCAGAGGCAACTGTACCCAAACAATTCTTTCCCTTTGCATCAAGAAAGTTTGTTGTTTATCTGA
GAGCTTAGTACTGTGCCTGGCACATATAAGGTGCTCAATAAACTTTTTAAAACCAATGCATCTGAGTGGCTTTATAATT
CAGCAGTTACTGTATGAACTGACTTACTATGTAGAAGAAAAAGTATTAGTTCAAAAAGAGGAGATTAAAGAATTCTTC
TTACATATATAAAACATGTCCTGTTCAGTAGCTTCTCTAATATTTTGTGGATGATTGGAATCCCTTTTCACTCATATTT
```

Fig. 6.93

AAGGATGTCATTATGTATGGGGACATTTGCTGTTTCAATACATAGATTTATATTTGCAGGGAACTTAAAGTCCCCGGAA
ATTTTTCTGAGATATTATCTGATTAATCCCCATAACAACGCTGTGAGGTGGGTTGTGGAGATATGAGGGGAAAAAAGGG
AAGGGATTTTCCTAATACCATACAGTTTGTGTATATGACGTAGCAATGAAAGTAGAACTTATTTCATCTGATGACCAGT
TCAAGACATTTTCTGCTATATAAATCTTTAATCTCTAGGATAGAAACTGTCTTAATTCCCTTTGCATTGACACAGCAAA
ATGTATATAGGTGGTCCATCTAATTCCATCTCTGAATATTCCAGCACTATCTGTCTTATCCACCTCCTCTATTCATTTT
TATTTCTAAAGTGTTCTCCCTGCAGGAGCTGACTCTTCCTTCTTTCTAATTAACTTTATGTGGAAACTCCTATCGTAAC
GAAATTGCACGTGTGAGTGTTCTCTCGCCTGGCTTTCTTTTTAGTGCACCTGGAGGGAGAAAGCTTTTCTGCAGTCTCA
CCACATTATCTTTGCTGAGAGAGAAACTTGCAGGAACTAAGATAACTGCCTTCCTGAGAGTCAACCTTTTCATCAAACA
TTCTTTAGTTTAGGCACAGCTTTGATTTTTGCAGAGGTTACCCACTTGTTCATAATTAAAATGCACTGGTTCATGCTATT
GGCAGCAAAGCACTATGGAAGAGTAGACAAAAACGTGGATTTTGAGGCAGACAAATCAGGTGTGAATACTGGTTCTGCC
ATTCCTGGGCAATTTACTTCATTCGTTGCTGCTGATTTCCTCGTTTGTGCCATGGGAAAACATATTAACAATGTCTACC
TCACAGGAGTATTGGAAAGTTTAATGATACATTATAGAGATTTTTTTAAGGATAAATGAGGTAGCAGAACATGCAATCC
TCTTCATAGACTGCGAGAGCACATCTACCATTTCCCTCCATCTCCACTCTCCTCCCTGCCCTACCTTGGGGGGCTTTGA
ACCCTGGAAGGCTGACCAGTCAGATGTACTTCTTCCTCCTTTGCCCTCCTGCTTGCTTGCATTTGGTTAGTGGGGAGTC
CTAGGAGGAGACAGATGGAGAAAGGAAAATGAGACTGCGTGCAACTGGTTGTGTCTTCTTCTTTTTAGGGTCTAGCTGTGT
CCCAGAGAGCAACTTCCCTTTTCAAGGCAGCCCACTCTGTGTGATGCTTTTTCCTAGGTATGGGCAACCCATCCCTCCT
AGGGTGAAAACTTCGCTGTTGCTAGTTCCAGGTACTGTGCCATCCTTTGTGGATTCCTCTACCCTACCAACATCTCCTT
AAAGAGTCTCTTTGTTGAACCTGCTGTAAATGATCCTAATAAAAGTATGCTGTCTGTTTATTCTTGGACCTTCAACTGG
TAGATTATTCATACTGGTTAGTGGAAAGTAGATCTGCCTACATATGTATTATTTGAGAGAGGTTAGCACTTATAGAAGA
AAAAACAAGAATGGGCTGTTTTTACTTGCCATTTGATACTAAGAGAAAAAGAAGGTAGTGATGATGGATGATGATAAAG
ATATGATGATGGTGAGACCCAAATACCTCCTATGAGCCAAACATTCTCCTAGACAATACTTTTTACCTCATTCATTTTG
AAAACACATATGTGTATTATTATCTTTCTTTTGTAGATCTGGAAAAAGTTCAGATAAATCAAATAATGTACCTAAAAGC
TACATAGAATGCCAGTGGAGAGCTGAAATTCCAACCTATACCTATTTGGTTCCAAAGTCTATACCCTTTTACTGACGCT
AGGTGCCTCCTAGCTATACAGGGTGAGAAGAGCCATTCCATCAGCAGTCAGTTTTCAACTGAATCTCTCTGCCTATTGA
AATTTCTCTAAGTTCTAAATTCCCATAGGAAAGTATCTCTTAATGATGGTCTTTAAATGATTTCAAGGCAAATTTTTTA
AAAACCTGGTTAATTCAGCAAAGCTTATCAGGTCAAATCCATTATTTGTCTGATTTGACTGATTTGTTACCATTGAGTC
ACTAGCCCAGTAGGGCAACTATTCCATGGTTGTCCCTAAGGCTACTCATTAAATCCTGGATGAATAATTAAATATTTTG
AATAAGTTTTTCTCTGATAATATATGTTTCCTACGGCTGTTATCTAAAGTTTTTTCTCCCTAGATATGGAATATTTCAT
TCAGTTTGTATTAATTTCTGTCCAATTCCTAAATTACATGAGTAACATAATTCTGCATTTTCTGGGACCTATAGGATGC
TAATTTGTAAAGGTGATTCAATTCCTGGAGGTGTACTAGCTGAGAACTTTCCATTGTGGATCAGCTCCTCCCTTCAAAT
CCTACTCCTTTAGAAAAAATCCATACACACTCAGAGAAACAGTATTTATCTTAGCAACTCACATTTGATTGTGCATTTT
TCTTTAATCTTCAGGCAAGCATTTCTATCAACCTTGGAAGAAGGCTTTGTCCCTTGTTTCCCTTTGATTGTGCATTTT
CATCCAGAGATATTCTTAGCCACAGTGAACCTTACTTCTCTGTTTCTACCTCTACTTTGCTGTTACGGGACCTCTTACC
TCCCGCAAAGTGTCTTCCTTTTATTTTGAGAAGACTTGAGAGGGTGACTCACATATATTCCAAACAAGTATTTTCAGCC
TTTAAAAAGGCTGTGTTCCTTGCAGGCTTCTCTGCTTTCATTTTGTATGTTTTTTTAAAAAATGATACATAGTTACTTT
GTTTTTTATCTTTTAAATGTATAATTCATACTACTTTGTACTTTAATATTGTCAATCATTTTAGCAAAACCAGCTCCTT
CAGACCTTAATCACTGTTACTCTTTTCCTTAAGTCTCAGACACATGTTTTTTGAGAAGCTTACAACAAACCCAAATGAT
AGAACTACATGCTGCTGTTAGCATCAGCCTACACCTACACTATTAGCCTAAACCTGCAATATCAGAGTTTTTGTGGTTT
GACCAAGCCTTTTATCCTCTTTCCCTTACAGTAAGTTCTTTCCCAGAAAGAAAATATCAGCCCACATGCAAATATATCT
TATAAATATGTAGGTTGTGTCCTGATGTAGCAGAATATCATAAATGAACACAGCACATATAGCTATTCGATTTGTTCTT
TCTTCAAAACTCAAAATGGAGAATCTCCACTGTAGTGTCTAGGAGAAAGCAAAGATTAGTTAGGTAATGAGGATGTCAA
TGTCATTCAAGTGGCAATTACTCTGGAATATTCCTTGAAATAACAGTTACTGCTTACTTAACAGTTATTGCTTATACTT
TTTGTTCTCATCTTCTCAAGTATTCCTTACAACCACTAAGAATAACAGTTACTGCTTACTTAACAGTTATTGCTTATACTT
TCATTGGATTGGCTTCAAGGTACTATTGATTATTGTCAGTGAAACAAGACGCAATGATGCAGTTGCTCAGAGGGCTCTT
TTCTTCACATGTAAGTAAGATTCCTCCAGCAGTGGATCACCTTAGTGATCCCTTAGTGAAAAATTGTCGAATCCTTAGC
TGTCCCAACAAGAATCAACATATACACAATTCAGTTTGCATCTTCATTTTATACATGTAACTTTAGGTTATGGCTATCA
TATCTGTTTTTTTTTTTTTTCAGCCACTAAAACTGTAGAGTTGAATATTTAATGGAAAACAGATGGTGCTTGAAATCTC
TAACTTACATTTAACAAAAGCTGTTTTGGAAATGTTGCTTATGTATACTTTTTTATTGCTTTAAACACAGCACCGTTTT
AGCCTGTTCCGATGAATTATTTAAGAATTAACTGTCCCAGTCTAAGACAGCATTTCAAAGTGCAAGTGTTAATCATAAC
TTGATTAAACATTTCCTTCCTTTTCTTTTCAGCATTCCAGTTGGCTTTTGAGTGGATACGTGCAGTGAGATCATTGACA
CTGGAAACACTAGTTCCCATTTTAATTACTTAAAAACACCACGATGAAAAGAAATACCTGTGATTTGCTTTCTCGGAGCA
AAAGTGTAAGTAACTTTTGTTTTCATCTATTTTCTAAACACATGTACATATAACATTTTAGTTTTGGTTTTGGATTTTA
ATGCTATGCTATCATGATTAGGCTTGTGGGAAACGTTTAGTCAACTTTCAGTTCTCTGACTGTACACAGCTTATTAACA
AACAAAGACTAGTTTGTCTGTGTGACCTGTTAGCATTTGATGAGAATTCTTTCAGCTAGTTTATATCTTACCATTCATT
CATATACGTCTACAAGCCTGATGTTGAACACTCTAATTTTTGAGTGCATGCAGAAAATATATGATTAAAAGTCAAAATA
TAGGGAATTTATGTCTTGAATATGAAACCTTCTGGGCAAGTCTAAAAGCATAAATTATACTTATGTTATTAGTTACTTC
AACCAGTCACATTCCTGAAAGTTCTTCCCTTCTATGAGCTTTCTACCCTGGCATATATCTCTAATTTCTTCTTTTAATCT
TTTTTTAAAAACATTTTAAATTGAACATCCTCAGGGCTCTACTGAAGGTTAAACCTTATTTCCAATTATGTAGTGTTCT
TAAGGTACCACAGTCATGATACTTTTTTAAATTTATATTAGTGTATGGAAAAAAGCAAATAGAGCAACTCAAAGAAAGCC
ACACAAGGAAAGGATGAAAGCAGACTATTAATTTCCATCAAGCAAGCGCTAAGAACACATCCCTTTGTTCATTTACCCA

Fig. 6.91

AGGGGGATAAGGCATCCAAACAGATGTACTTGTGACGTAGGAACATTAATTTGAAGGCATCAGAAAAACCACAAATGCA
AACACATTTCTCTAGTATGGGAACACTTTGTGTTATAACCAGTGATTCTATTGTGGTGGCCCAAGGCCATCTTTCATTC
TTTCTATGGAGTTACCATCCAGCTTTTAAGGGTGAGGCATGAGTATGTGCAGATAAAGTATAGTATGCCCAACTGTGTT
CAACAAACTCTGAGGAATTGTTCAAATTTATGTGTTTTACTGGTTGGTGTTTGGTTGTATGGATTATTGCATAAAAGAA
GGGGCCATGGATTTGAGCCTGGAATTCCTTTTATCATCATACAAAGTCACTCACTGTAATAGGAGGTGACCTGCATACA
AATGTTTTCTGTTATTGAATTTCAGAGACATGAGATCTGAAAAAAAAAATGTGTAAAGGCAAAAGGGAGGATATTTTGAA
TAATTACAAATGTCATTAAACATTTCCCTATTCTTGGAGGAAAACATTGTAAAAGCAAATGATTAACTGAGCGGTGACT
TTAAGGAACTGAGACTACTTAATGATGCTAGGAGACTTCCTATTTGTATTTGTTTAATGCAAAAAATTTTATCTTGGTG
GAAAGGCTCAAGCTTTCCAGATTAAGAGAACCTGAGTTGCCTACATTTGTCAAAATGTAAACAGTAGAACTCTCATTTC
ACTGCTTCCTAGCTTCTTGAGAATCCACTGGCATAAAGAAAATGCTTCCACTTAATTAAATGACTCTAAGCAGAAGTTA
ATGTTTATCAGAAAAAAAGAGCCAGACTTATTGCCTAGTTAGAAGTTGTCACTTTAGGGCTATAAAATTTTATTTTGCT
CTGGTCTGAGCATATAACCCTCCAACGCTTACGTTTTTGCCATAATATAATCCAAAATTGTATACTTAGAGGTAGATTT
TTCCTGGTTTTGATGAAGAAAGTTTAAAGAATTAGTTCTCTTTAGAATCTGAGGGATGTTTATCTTTGGCATTCTGACAT
·TTGGAAATCCAAAGAAAATGTTCGAAATTTAAAACAGTCTCTGTTCCTCCCTACATGCCTCTCTTGTAGGTCTTGCATC
TTTCATTGTGAAAATTTAAGGCATACAAAAGAGGTAGAAAAAAGATCATCCTTAAACGATCACCTTACTTAGTGAGTTC
CTCTCATGAGAGGGGTTCCAGTGTGTCTTGGAAGACCATTTAGTCACTCTTCAACTCAAACAATTCAGGCATAAGATGG
GTGGTTAAACTATGTGAGTGTTCTGTTTCCTACCAGTTATGAATTTCTATGATTCTATACCATGTTGTGCTCATTCGTA
AGTTGAATCAAAGACCAGTTCCCAAATAAATATAAATCAAGGCATCAGGGCAAACAGAGTATATTAATCAGCTTGGGC
TACAATATACCATAGACTAGGTGGCTTAAACAGAAAGTATTTTATCACAATTCTGCAGGCTGGGAAGTTCAAGATCAA
AATGCCAACCAATTTTGTTCCTGGTAAGCTCTCTCGTCCTGGTTTGCTGATGGCTTGCCGAAAGCTACTTTTTCCCTGG
GTCCTCACATGGCAGAGAGAGAAAGCAAGATCTCTGCTGCTGCTTCTTTTAAGGTCACTAATCCATCATTAGGGCCCC
ACCTTCATGAATTCATCTAACACTAATTACCTCCCAAAGACCCCACCTCCAAAAACCATCACATTGGGGGGTTAGGGCTT
CAACAAATGTATTTCAGGGGGACACAAACATTTGGTCCATAAGAGAGTTTGATATTCTTTTTTTTTTTTTTTTTCTGAGA
CGGAGTCTTGCTCTGTTGCCTAGGCTGGAGCGCAGTGGCGCGATCTCGGCTCACTGCAAGCTCCGCCTCCCTGGTTCAC
GCCATTCTCCTGCCTCAGCCTCCCGAGTAGCAGGGACTACACGTGCCCGCCACCACGCTCGGCTAATTTTTTTGGCATT
TTTAGTAGAGACAGGGTTTCACCATGTTAGCCAGGATGGTCTCAATCTCCCGACCTCGTGATCCGCCCGCCTCGGCTTC
CCAAAGTGCTGGGATTACAGGCATGAGCCACGGCGCTGGAGTGCAATGCCGGGATCTCAGCTCACTGCAACCTCTGCCT
CCCAAGTTCAAGCCATTCTCCTGTCTCAGCCTCCTGAGTAGCTGGGATTAGAGGCATGCGCCATCACACCTGGCTAATT
TTGTCTTATTAGTAGAGACAGGGTTTCACCATGTTGGTCAGGCTGGTCTCAGGTGAATTTGATATTCTTAAGGGATGAT
TGATTTAATAAGTCACTGTCTTGTTTAAGCCCAAAGGGTAGTGACTAGTATAATGGAATCTGTATGTTTTCCCAATTTG
GTAACACTGAAAATGATCTGGTCAACATCTTTCTCTTCATTTCTTATTTTCTAAATTTTATGTTAGGGACATTCTTACC
AATGATTTTTGAAGTCTAATAATCACTTCATAGACCTAGAAGCTAATTTTAATTTTACTGAAAATGATTTTCCCCTTTC
TTCACTTATTCTTAGTTTGCTTATTTTATATTGTTCTTTCTTAACATCTAATCCAATGACAGGTCTCTGAAGTATCTTGT
CCTATCAATGGATTTACTATTTAACTTTTCAGATGTTTATATATTTCAGAACTGACTATCTCAGTTACCCTTCTTCCCT
CTGCTAATGCCTGCTTCTTACTGGCTTTTAGCTAATCAGGGGAGGGAAGGAAAGGTACAGACAAGAGAACAGGATATA
AAAATTTTTGTTGTAATTATATATTTTAAAATCAGGAGTTAAAGTAAAAATTTTAGCATTTTCTTTTCTTTATAATATC
CCTGTTATTCTCCAAACTCATGAAGTTTTATAAATCTTCTCATCTGGAGAATAAACTAGATAAATTTTAATTCTTATTC
ATTAATAAATTTTCTTGTGTTATTAAATTCTAAACTTTCTTGTAGTCTTTTGCATTAAAATTCCCAATTTTCTAGTACACT
GAATCATTTTCTTCCTACCAGTTTGAAGATATTCAAGATGTCATCTACTTAATTGGATTTGTTAATTTTCTTGATGAAA
TTGCCCTTTATTTGACTTAGGATTCTTATGGATGTTCTACTTTTGGATAAATTTGTCACTCACCAGATGTCTTTGATGT
TCCTTGTTCTTTTTTGATAGAATTGTTGTTCAGTGTTCATTCAAAACTGGAAGCAACTCCTTGGTGTTTGGAAGACATA
GGGTATCTGGCACATTCTTTCATAATGAATTTAAAGCCCACCTAAATAGGCTGTCAGTTTTGAGTCTGGGGGCTAGGAG
CCCATCTGGTGAAATTTTTCCCAGTAGGCTCCCTAGAATATTGCTTATTTTCATGGAAGACTTTAAAAAATTGTACAAA
CTTGGTAAACTAAAGTGATTTTATTCTTAAGAAAATATATATTTTTTTCCATCTGTACCTTTACATTGCTGAGTTTTTA
GAAGCAGTCACTCACCTAATGACCCAGGGTGAGTTATTTATGCCATTTTATCTGCATTTTTTAGTGTCTTCCTGGAGTG
GGCAGCCAAAAAATAAATTTAGAAAAGATGGTACATTTTGTTATGGCACTTCAGAGCTATCTGCATTCATTTTTCCTCA
CAATTACCCTGTAAGATAGAGTATTTTTTATTTGTATGGTTAAAGTAAAAATAACAAATTCCTTCCTAGATGGGCTTCA
ATTCTCTACTATATTCAACCCCACCTAAGCTACGAGATTGACAAAGCTGTTGCAGCTTAGAATCCTAATAAAGTTAAGG
ATGTATAAACTGATTTTTTTCACTGGTATTCACTCAGCAGGGTCCAAGAAGGTCACGGGAGATGCTGTAGGTAAAATCC
TTTGAAAAGCACAAATACAAAACAAAACCAAGTTCCTGCTTGTGTCTTATTCACAACAAATTGATATGAGAAGTTGT
AGTTTTAGAGATTTCACTCCTTTGGTCCTTATACCTGCCTTCCAGCTTGTTCTCCTTCTTTACAAAGTTCTACAAATAT
TTACTGAATGGCTACTGTGTGTTAGGCTCTGAGTATCCAATGTAAGATACATGTTCTCTCTACCCTCATAGAGCATATC
TGATAACTAAATTTATAATTTTTACAATTTGTTAAAATTATTTCACCACCCTGTCTCCCTAACCCTGATCACCTAACTG
TATCCCATTTTAATCTTTAATTCTTTTTAGGAAGGCTGAAGTTCTGTATCTCTCTGAAGTCTCTCTGTATTTGCACTTC
ATCACGTTGTGGGCACCCTGTTTAGTAGGGATATTTTGAGCTGTGAATAATGTGCTATAATTGACAATTATGCTCATTG
TTGCAGAAACATAGATTATTTCAGAGTAGATAAAAAGAATTTTTCTCAGATAATCAACAATAGTATAAGAATAGCCAAG
AGTATAAATAAATGTGAGAATGACAGGAAAGGTCTGAAAGAACAGCGACTGATCAGCAACCATTTGCTTTAAATCAATC
ACATGTCACATGACTGTTTTAGTGGCCAAAGCCAATCATTAAATTGTCATCTTGGAAAAAGTTCCACTTTTTTTCTGTAC

Fig. 6.95

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TGTGATGTGTTTGCTATTTCATTTTTGGACAATTTCAAGTTGAAGTTGAATTACAAATGATTACGTCCAAGAAATAATT
GCATATATTGTACTTACCAGTGTAAACACCACAGATACAAAAAAATGGAGAGAAATCTTGTTGCATTGTTAATGAATAT
TAAAAAATTATTCTTACAGACAAAATCTCTCTA..ACAAAGTGAAATTTTGAAGAAAAGCATAACTGGTGGAGTTTTCT
ATCAGTAGTATTTGCTCTAGAAAAATGTTTATAATGGCTGCTTATTTTTAACTGAAGGTAATTTTCTTTTAAAATTTTG
TTTAGCTTTTTTTTCACTATTGAAATGGCAAATGTTTTGAAAACTAAAAAAATTGACTTCAATTAAAAATTATTAGTGTGTT
TACATTGTTTCCCATGGTCAGCACTTTCAAGGGCAGAACTGGAATTGTCCCTGAGTTATAGTTCTGGCTTTGCATCTGT
GCATCTATGTCTTGGTAGAAGAAGTGGGAGTAAGAGAACCATAAGTAATTAGTTTTATTATCAAATGCTTCCAATAGCT
TGCATAATTTTTTTCAAGGGTAAAATGAGAAAAACAGCATTGAAATCACATTAAAAAATAAAGCAAACAAAAAATCCAA
ACCAAATCATCAAGCAGAAATCATTTTAAAAATGCTATACAATAATCAATTGTGTTCAATACTTCTGCGCCAAAACCTT
TAAGTCTTAAAGCTTTCCTAAATGGAATTAGTCCATCAATAAGCAGATAACAACTTTACTTTTATTGTTGAAAAATGTC
ACTCTTGTCAGTTCACAAGTTTCCCTAACTTGTGCAAATTAAAAGCCATGACAGCAAGGCCAAGAATGGTTCGTTGGTA
TATAGTTAGAGAGACCCTGACCCTGCTTTCTTGTTCTTCCTATTACCCAGGATATGTCATACGCCACACACATGGGGGT
AGGGATACCGCCCTACTTATAGCATAAGAACTGAGAATAAGACATTTAGCTTATAGTCATACGCCACACACATGGGGGT
CCCACTGCCATTCTTCCTCAGGTAGTCTGGACTTCTGTTTTGGCAATATTGCTTCCGTGAGAAGGATTTGACTGTA
CTTGCAGCCCTCAACATCCTGAATTTAATACAATCTACGATATTTGTTAAGCCCTCACCAACTATTTGACCAAACTATG
ACAGTTACATCCCACGGAACTGTATACCAAATGGTACCCCAGCAGTTCAGAGAGTTAACACACCACCAATTCACTTCCA
GGTTGGGCTTTAAGATTTTCAGAGGATCTTGCTTATGAAATCATATTGGCTTTTAGATAAATGTATTAAAATTACTGAA
AACAAATGATAACACCACAGTACCACAACCTAAACAATCCTTTCTATAAATTAATTTAAAAACAAGAAACAACAAGCTTT
AAATGTCATCATCTCTGCTTATTGTATGTCCTTAAAAATATTATTTAAATGTCCAACTTTTATTTTTCTAGAAGAGGTCA
TTATATAGCATTGATTTGCCAGCAGGGTTCTATTTGAACATACCAAGAGACCTAGACATTGCTCAGAAACAGTAGTCTC
AAAATAAACAAGGGATTGGAGGAAAAGATGAGAGATCTCCAGTATGTCTGCATATAAGGGCTGAAAAAGTAAAAGTTTC
CAATTGTTTTTTTTTTCTTTGCAAGTGCTCATGCAGGGATTCAGAACATGCATTCCTCACCCAACATAAATGAAATAATT
GGCAAGTAGTCAAAGAAGACCATATCCTTGAGTGGGTAGTATTGTTGTCTATTTGGAAAGCATTTACATGTTTTGATTT
CCTGAAACAATGCTGAAAATGTCCTAATGCAGGAAGGGAGAATTGAAAACAACCACCATAAAATGCAATTAGATGTTGG
TAAAAGTGACTCAGAATAGACGTAAGTTAAAACTTTCTGACAGGGTTTTCTAGCACTGGGACAGTATTTTTTGAGGAAA
TTATAAAATGGTCTTATTTCAAGATCTTTTAACAATCTGATGAAATATTTATGGTTCTTTAAATTTGTATTGTTAGTAT
CATACTCTGAATTATAAAACATTTAAACATTTTAGTTTATAATCTTTAACTTCTCACTATTATATTTGTATTGTTAGTAT
ATAATTTGTGCATATGTAGACATTCATGGAGGAAGATGAACATATATGTTAATTGGCATCTGCTCATATTTATAAATGTGTAT
TGTATACTTTCATTACAGTAATACACGTGCATTCATTAAATTATCTTGTGGCTTAGCTTTACAAATTCATTTAAAACTAAAGT
GACTTTGGGTGAATGACCTCACCGGTGTCTATATCATTAAATTATCTTGTGGCTTAGCTTTACAAATTCTTACCGTTACAT
TTTCTTTCAACTCCAAAGCTATGACAATGTATTTTCAAGATTGTGTACTTCTTAGAACAGGCTCAATAATAATTTTTCA
CATTATGAACTTTGCAGTCAAAGAATAGGTTCTTCTTAACCTAACAAATGACTATCCTTTCCACCCAAAGTATAAACAG
GATCCTGATTAATAAATTGAGTTCAAAGAATCTCTGCCAAGTATGCAAATCACTGCCTCTGTTTGTCGGACATTCATT
GCCTCTAAGAATTGATGGAATTGAAAATAACCTCATTTTACTGGGACCTCAGAGAATTAATTATTTAAATTTTTTGCTG
CTTTAAACATTAATTTTCTTAATTTACCCATATATGTTGCTGATAAGAGCTGTAATATTTTGAATGGTTGTGCTTTGAA
GAAATCTGAATCCTTTTGCTTTGTATTCCAATGACAGCAGCTTTGACCAGCGACCAGCTCTCTTCTGAAAACTACCATT
TTGACCCCTGCATTTCACCTGTTGTCCCACCAAGCCTGTCTGTTTTTCCATTAACAGAGACCTCTGAATTGTTGTGCCA
AATGTGGACAGTGTTGTTTCTTCCTCACTTTCCTGAATTATAACCAGTTCCAGGCGGTAACATGCAACCGAACTTTACT
GAGTCCTGGACATAATGGGAAAAAATCATAGTGTAGAAAGAAAACTATAGGACTGAATATAAAAAAATAATCATTTCTG
GCATTACAGGACAAACCCAGTCCTTTCTGCTTAGTTACTGACCTACCCCCTGTTGCTTTGCTTATCTTCCCACAGTGAA
ATGTCTTTCTTCTTATATCCTACATGGTTTCCAGGCCCTTTACTCCAGGAAAGCCAGGAGAAACGCCTTATTCCAAGTT
CAAGTAAACATAATAATTTACAAAGATACAACTCTGCCCACAACAAAAAACTCCTTTTACAGCGTTATGCAAAGGCATT
TAGACTGGAACATCTATGTTCCAGACACAGACCTTAACCAGTCTTTTGTCAAACTAAAAGAGCAATCTTTCCTCAAAGC
TGGAATAACACCTTTTCTTTTAAAATAACATTTCGTGTCTCACACTCCCAGATGTTTTCATTTAAGACTTTAGAAAATA
CTGGGATCAGTTATCAGCCAAGAGTACCCCCATTCAATAAAAATATTTAAAGACATGGAAAAATCAATGAATCCAAAC
AATCATCATCCTCACCAAACCCTTATCATTTCTATAACTCACAGTAAATAATCTCAAGTTCTTTATTTTGGTAAATTAA
GAAATTCCAGAGTAAACTCTCTAGCTTCTGATTTAAGCTCAGAGATGCAGAGAGCTTCAGAGTGTCGTTCTCATTCTTA
CCATAAGGAAAAATCTATACAAACTACCTATTCATTAATTTAAAAAAAAAGCACCAGCAAATTTAGGTCGTAGGGCAACC
AAACAACTCCAAATCTGGAAAGAGACAGGCACCTGCAAGAAGAAAGAGGATGGCATCATTTGTTTTTCCTTGGGTAGAC
ACCACCAGATGTCATGTAAACCAGCAAGATGATTCAGCTAAACATTTTAATGAATTGCTAAGGCTGAGTATGGGCTAGC
ATGAAAATGTGACACACTGGAGGTGGCAGATATAGGGAGTGCATCCTATAGCAGGCTTTTCCTCCACAAACCCCACCAG
GCACTCACAGGAAAGACTGGGGAGAACAGCAGCCACCTTCAAACCCAACCATTTCCCAGTGGAACAAAAGAGTTAAT
TGGCAAAGAGAATAGCACAAAATCATTGTCTTAGGGAACTGGAGGAAACCCATTGGTGATGGTGGCAGTAGGAAGAATGA
TCGTGGTGAGGGGAAGAGAAAAAGTAATGCTCTATCCCCAGGGGTGGGGTATGGAATATATGCTAGGATTTGCACAAC
AATTGGAGAAAGTTGCAGGAGCACTTGTGAAGGCCACATTGCTGATACAGAGGGTACACATTACCTACCTAAGACTGAGT
CTTAATCAGAACATCAAGGAATGCCCTTCCTCCCTGGCCTGCTACCAACAGCCTAACAAGTGTTGAGTAAAAATAATAT
GGGAATATGGTTGAATATGGAAGAAATTCAAGAGACACATTCTCTTTAGGGCCCAGCATTAAGGGAAGACCCAAAGCTA
AAGGGGGAGCAAATATTAAGAAAAATAACAACTGGCAAGCCATTTCACAATCTATTCCTCTTTAAGAATCCAAAGTATC
TATCTCAGTATCTACTGTCCTACACAAGATATCCGGCTTTCAGCAAAATATTATGACCATATGAAAAGGCAAGAGAAAG
CACTCCGAAGAGATAATACACATAAACATGTGATATATGGGACACATATAAAAATTATCACACAAGGAATTTAAAGTAA
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Fig. 6.96

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CTATGATTAATATTTTAAAGGTTCTAAAGGAAAAGGTTGACCACATGTGAGATCAGATAGGTAATTTCAGAGGAGATAT
GGAAACTAAGAAAAATCAAATGGAAATGCTAGAAATTAAAAACATAGTCAAAGAGATTAAAAAATGTCTTTGGGCTTGT
TAGTAGACGTGAGACAGCTAAGGAAATAATCACTGGACTTGAAAATAAGTCAGTAGAAATTACCCAAACAGAAAAAAGA
GTGGAAATAAAAGGGAGAAAGAAAGAACATAACAGAGCATCCAAGAACTCTGGTACAATATTAAGTGGCATGACATATC
CATAATTGAAATGACAGAAGGAGAGGAAAGAGAAAAAGAGGCAGAAGAAATGTTTGAAGAAAAAAATCGCTGAAAAAAT
TTCCAAAATTGATGATGGACACCAAATCACAGACTCAAAATCTCCAAGTATATCAAGCAAGATATATACCAAAATAAAC
ACACACGTGCAGAGTCATGTTATATTGAAATTGCTAAAAATCAACAACATAGAGAAAACCTTAAAGGCAACCAGAGGGA
GGGAATGATAAATTACCTAAAGAGAAGTAAGGATAAGAATTACAGCAAATTTCTCCTCAGAAATGTTGTGAGCAAGAAG
AAAAAGGAATGACATCTTTAAAGTATTGAAAGAAAAACAAAACCCAGTAACCCAGAATTCTATGCCCAATAATAATACA
GTTAAAACAGGAATTTAAAATTTTTCTGACTGAACTTTTTAGGTTCCTATTCTGTCTGATGTCCCCAAAGTAATATGCAC
CTATATCCTCATTTTATTTTGTCTTTCCTTTTTCAAATCTGAGAGGGTAAAGCTGTTTCTTCTATTCAAGACCAACACC
TCTATTGCCTCAGAGGGTTAGTCAATTGCCTCCTCCCTCTATAGGCTCAATTGCAATTTAGCTAATCTCTCTTCTATTT
CTCCTTTTTACCTTAGTTTATGAACTAGACCATCCAGTCCTTTCTGTCACTCCCGTCCATGTATATCTCTTTTGACAAT
TCTAGTATTTATATTTTTCTTTCTAGTAGTTATTGCTATTTCCTTCTGAGACTGCATGTTCCTTAAACAATGCTGTTCC
CCAAGATTCAGTCATTGGCCCATGTAATATTTTAAGGTATGAAGACACCTGAAAGATCATAAGATATCTCCTTTCATGT
CTTAACTATCTTCTATTTGATTATCTCTCAATTTCCATATTTAGTTCCCATTTCTACTTTTAGTCACAGAACTCTAACC
AAGTGCTTGCTGGACATTCATAACATCTACACAACATTTCCCAAACTAGATGCATCCATTATTCTTCCTCATCTTTAAA
CATACTGCTACATATCCTAAATATTTTCTATTATATTTACTGCCTCAGCTATCCAGTTGCCTAAGGCAGAGCCCTGGGA
ACCTTCCTCAGCTCCAGACTCATCATCCCTCCCAACATCTAAACAAGCAACAAGTGTTACTGCATGTATTTGTGGGCAC
ATATTTCTGAGTCCTCATCTCAGCTTGACTTCCCTTTCCACCAATGCCCTGGGGATCACCTTATTTTTTCTCCCCAAGA
TGGTTCTCATTAAAATATACCACAGTAGTGATAAGTAGTAAAATTATATATGGAAGCTGCTGACTGTTGCTCCATTAGC
TGTTTCTGTGGCAACTCTATTTTTTCCTATTAATAGTGTACTAAGTGTTTTGAATCTATGTCTTTTCACAACTTCTTTT
GATTACAGTTTAAGATAGCCTTTAAATACTAAATTCATATTAAGGTGGATCTGATTGTGTGTGTGTGTGTGTAATTT
CTGTTACAGCCATCATTCATAATTCATTATTTGTGAGGAATAATTATAGCAAAGACTGATTAGATGAAATGTTAGATTA
TTTATTATACCAGATTCGAATGGAATTAAAGTCTAATATATGTGTTAATGTGACTAATAAGAATAACAATAATAATTTA
GAATTGCAAAATGCTTTTAAGTTTCCATATACTATTACAACATTTTCTTGTTTTTTTCCTTTATAAAAACTTACAGGTA
GGAAAGGCAGTTATATTTATCCTCATTTTATGAGCCTTAGACAGATTAGGTGAGTTAGCCAAGATGCACAACTAGTAAT
GGTGAATCCTAGATATACTTCAGTTCTTCTGATTCCAAATTCCTTACTTTGGACACTAAACCTTGGAGCCACACAAATG
TAACTGTAGGCATCACTTATGTACTAATGTACCTGTTACACCATGTGCCTACTGGAGAGAAACTGTAACATATTTATAG
TCAATTATCTATATTTATAGCTATCCTCACAGTTGGATAACAATGTTATATTGTAAACTCTAGATACTTTTGCACTCTA
TGACAGAAAGTGGATTTTTTGTTTAGGAAGACATTTTAATTATGAATGGAGGCTTGTTGGGGATCCCTATTAGCCTTTG
TTAAATCACAAGGAAGGTAAACTCTTTTGCATAATAAAGTATTAGGATTTTTAAACTTGTCCCCTGACTGCTCAAGAGA
TGAAGTTTGGGGAGCGGCAGTACTTTGGAGGACTCTGGTAAGTGAATAGATATAAGAAAGTTCTGCAGGGATAGAGCCA
CATGGCAGCAGACAGGGCTTAGGAAGAGGGACTACCAATTGTCCCTGTAGCAAAGATGGCCTGGTTAAAGCCTCTTTAG
AATGAATGAGTTCTAGTAACCAGTTTTAAAGATTTGTCTGTTTCTTTTCTCCCCTCAACCCTTTCTTTAAACTTTCTGT
AATCACAACAGCATTGTAACAATTTTGCCTCAAGCTGGGAAGAGAAGGGCTGTGTTCCTCCTAGAGACAAAACAGGGAA
GAAAGGTAGCAGGGGCAGAGTGAAAGCAATGATTGCTGTGGAATTGGAACAACTGTGCAGAGGAAGCTGTGACAAATAA
TTAGGGTTGGTGAAGCACCTGTCCCTTTGGAGGTATTTCCCAGAAATACTGGGAAGGGCTCTAAATGTCCCATTGTTAT
GTGGGGAGATGGTTTTGTTTATTGTATTGCCCACAGAAGTCTGAATACGTTCAAATATTTTGCATATTTGTGTGTGTGT
GTGTGTGTGTGTGTGTGTGTGTGTATTGAAGTACTTGGCATAAGGTTATGCTATTTCAAATTAACTTATAAAGTAATTT
GGTTGATGCAATATTCTGTGATCAAAATTTATTGAGGACCTCACCAAATGCCAACATGTGTGTGTATATATATATATTA
TATTATATATATAATATAATATATATATATTCACTCATTTAGTTCTCAAAACTATTCTATGACACTAATATTTTATGTA
ACTTTTCTATGATACGATTATTATTCCTATTTTTTGGATAAACAAACTAAAGTACACAAAGGAGCCAATACTTACGTCT
AAAATACCTAAAACATACATATTTGAAATAAGAGAATATGAGGAAAAAGTCTCAAAATTTTATGTAGGTTTTAATAATA
AAATTGAAAAGTCGAATCATTTTAAAAACCTTAAAGTTAGTATATGGAAATATTTGGCTCAAGAGTGAACTTTAGACCTC
TTGTCTGGATATGAGTATGGAATACATTATTCAGTTTCTTTTTTATAACTTTAAATAGCTTGTAAGAAGACACCTATGC
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GATTAGAGGATGGTCATTCAGATTTCACTGGATTTAAATCCACAGTGGGATAGGTTTTAATCCTTTCTGGAAAAAATAT
TTATAAAGCCTATGTGAGTTTCAGAATAGTAAGAGCTTTTATTCTGTCTCTCCTTGGATGTGTTAACCCTTCCTTCTAC
CTCAGATCCATGCATTTCTCTAAGTTTTATACTTTGTTTAAATTGAGCTTTCATTGTTTCTATGACCAATTTTCAATTT
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CAGAATTGTCTGAAGAGTTTATTGTAATTTAATAAGATGTTCTCTTCTGTTGTTTCATCTATTATGTCATTACATATCA
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```

Fig. 6.37

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ACAGAGGCCTCATTATGATAATACTATATAAAACATTTGTATGATGCTTTTCAGTTTACAAAGTCCTCTTAAAAAACAT
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TTACTTTACCAAGCTGAGCCTTATTTGACTCATCCCAAGTAATATAGGTGATAATGACTACATCATAGGATTGTTATGA
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TTAGACTCTAAAAAGGTCATGGTCAGTATAATTTATGAGGTAAAAATGGCATCAAAAAAGAAAGAAAAATGCAGAGTAC
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AATATAATTTGCAGGTCCACAAAAAGTAAAATGATATAAAATTATGCTAAAATCAAGAAGAATGGAAATTAATGGATTT
AAAAAATTATTGTTATTGCATTCTCTCAATTTTTTAAGACTCTGTCTCTGCAGTAATAAAGGGAACAGAGGGAAAAGTG
GGATTCCAAAATAGTTTATTTAGACTTGGAATGCTTGCATGCTTATATCTCACTCAACATGAATACGTCTTAACTAATG
GGGACTTAATGAAAACTTTATATTATTATTAATTGATTGTGGATTAATAAATGAATTAAAAATGGGGTTAGAAATAGTG
CGTTATTTCTTCATCTCTTGCCTTTATTCAGCTTTTTCTCTTTGTCCATTCTACAGTAGGAATGAATTCTATCTCATAGA
TAGATTTAGAAACCAACTCCAGCAAAAAAATTGTTGTGCTCTTTGTGAGGGAAGAGGTTAGATTAGGATGAATTAAATA
TTTCAGTTGAAAAAAATCTGAGCAAATTCATTTAAGATATTTTGAAAACTCCTAAACAGTATATAAATATATGAAAGTT
TAAACAATAATAAACAGTGGAGGTACATCTTTTGTGTTGTAAAAAATGTAGTTGATCTTTTTCTTTAATGTTTCCTTAT
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CTGTTAAAAGTAAAAGTGTCTTTTATAAGCCTCATGATAACTACAATGGAAAAATAACTTGTTAAAATTATGGAAACCT
CGTCTCTACTAAGAATACAAAAAATTAGACAGGCGTGGTGGCGGGCGCCTGTAGTCCCAGCTACTCAGAAGGCTGAGGC
AGGAGAATGGCGTGAACCCAGAAGGTGGAGCTTGCAGGGAGCTGAGATCGTGCCACTGCACTCCAGCCTGGGCGACAGA
GCAAGACTCAGTCTCAAATAAATAAATAAATAAATTTTATAAAGTGTTTTTTATAAGCCTCATGGTAACTACAAAGGAA
GACCCTATTATTGGTACACACAACACAAAATGCAAGGAATCAGAATACACTACTAGAGAAAATCACTTAACCACAAAGA
AGGGCAGTAAGAGAGAAATAAACAAAGACTCTACAAAACAACTAGAAAACAGTGAACAAAATAGCAGTAGTAAGTTCTT

Fig. 6.98

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ACCCATAAATAAGCCCTGAATGTAAATGGATTAAAATTCTGCAATCAAAAGACAGAAAAATAGATTTAAAAAAAAGACCC
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AAGCTAAATTAGCCTAAGCAGCGTGATATCAGCATAAAAACAAACAAACAGATAGACCAATGGAACAGAATACAGAGCC
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GGTGTGTACAGATTTCCAGAAAGTCAATGGGGATGTCCTGGGGCTAGTGGGCCAGGCTTCTTCTTTTCATTTGTATTCA
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TCTGCCACTAAAGATGTATGTGATTTCTAGAAAAATAATTTTACCTCATTGCATCTCAGTTTTCTCCTTTGTAAAATAG
GGATAGTGGTAAGTGAATGATTTCTGTGATCCTCTCTAATGCTAAATAGAATGAGAATGTGCGAAGCCTTTGTTATCTC
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CTCAACAACATCTGAAAGTAAGTGGTAAAATAGACTTCCTTTTATATAGTCTAAACTCTCAGTGGCCAGATTATTAGTT
TCTTTATTAATTCTGGATGGTGAGGAAGAAGGGGGACATGGGTGATAAAGTTAATGTAATGTAATATTGCAGATTGTAT
TATTAATGTAAATTCCCATTTGGAACTCAAAAGCCAAAATGGATCTGAAGTCAACTTATGCAGTCTACTTTTTCAGAAG
AACAATTAAATAGTATGAGGTAGAGACAACAAAATACCAGGTTTATGGAACACTAGAAAGTGGAAAGGAGCCATGAGAG
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Fig. 6.99

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TTATTGCGGCCCTTCTCTCAGTGTCTATTGATTTCCATTAGCATCTGGGGTACTTAACTATTTCCTTCTCCTTCTACTT
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GGACTAATAAAGAAAGCAACTGAATAAATACTTGAATAATAAATGATATGTGTTTTTCACACTCTGGTCCAGTTATTTT
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AAGACAGTGATGTTGATGATCCTTATCCTGTGTAGGCCTAGGCTAATGTATGGGTTTCTGTCTTAGTTTTTTAACAAAA
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TGTAAAATATGTTGGTGTTTTAAGCTGAGCATTATTACAAAAAGTCAAAAAGCTTAAAGAAATTAAAACGTTTATAAAA
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CGGTGGCTCACGCCTGTAATCCCAGCACTTTGGGAGGCCGAGGCGGGCGGATCACGAGGTCAGGAGATCCAGACCATTC
TGGCTAGCACGGTGAAACCCCGTCTCTACTAAAAATATATATTTAAAAAAATTAGCCAGGCGTGGTGGCGGGCGCCTGT
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ATCCTCACAATAACCTTATGAATTTAGTACTATCAATTAGCACCATTTTTTTCCATGGGAAATAGGATATAGGATATTC
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CACCATTAGTTGAAATCATTGCACTATCTGAGAAGAAAGACCATGAAATCTAGGAAGTTTAAGAAGTTAGTTTTCGTTA
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TCTTGTACCTAGAACACTGCATGGTACTTAGTAGATGTTCAATAAATATTTGAATGAATGAATGTATAAGTAAAATAAG
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CAACTCTCTTTCATGGTATATTATATAAGCATCATGAGTTAGGTTTACTAAGAGCTTTTTTAATATGAAAAAAATGCTA
GAAATTGTGCCAAAATATTAACAGAAACTAGGGCAGGTAAGATTATGGGTGATTTGTTTTATACTTCTCTATACTTTT
TGATAATCCAGAAGTGATTATGTTTATACAATAGACTATGGCTTTATATGAAGAAATGAATATAGTCTAGTATTGTTT
TTATTATCTAGGAATATACATGTAACTGAAGAATTTATGAGTAAAGTTTAATATAAGCAAGTAACTGGGACTTCTGGAG
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Fig. 6.105
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GAAAGCTCACTCTAGGGGAAGCTCTCAAATGTGTTTTCAATCTTGGATTCCAGTCAAAAAAAGACATGAGTTACTTGGA
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CCCTAGTAACTCTGAGGACTGTACCCTTGATTCTTGGATTGTTAGACACATTTCCACTTCTGGCTTTGGCCCTGCAACC
CTGGTACCTAAATTGAGCAGCTTGAGCACACAAGTTCCTCATTCTTTCATTATGTATGTCCTTGTTAAGAGACTGTGAA
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AAAGCTATGTGAACTATTGTAATAGGTTCAAGAACACACCTTAATTCTTGGACTGTTTTTTAGACCTTTAGAAGTCGAA
GGCCTCTGAAATTATATGAGCTTGGGTGTTGAGATAGAGGAAACATGAAAATTTATTCTACCTGAACCTCTGCAGTTAT
AAACTAGCATCGTATAATTTATTTATTTATTAAGTCAAGGTTTATAAGTCATTTCCACAAAGCCTAGCACACAGTGTCT
AGCATATATAGTAGATGCTTGATAAATATCTGTCAAGTGAATTTCAAATACTTAAATTTTGTTGTTAATACATTCATGT
AATGGAGCCATATCATCTCATATGAATGATTGAAGAATACAAAAACTCCAGACTTGAATGCAGAATATATAAAGAACTA
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AAGAAATCCAAATAGTCAATAAGCATATAAAAAGGGGCTCAATCTAATCATTGAGGAAAACCAAACCCATAATTCAATA
ACCCTCAATCACTTCCTGTGATAGTGTGTATTGGTAATTGGCTTGAAAAACTCTTTGGCAGTATCTACTAAAGCTGAAC
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GGTGCTGCTTATGTGGTGTACTCTTTATAAAAATGTATTGATGTTTACACTTACGTGCAATTTTTGTATACATATTATT
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CAACAGCGTACTAAAAATACTAAAAGGATGATATACTATGATCAAGTGAGATTTATCCCTGGAATGCAAAGATGGTTCA
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Fig. 6.101
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TGTTGGTAAATAATGAATGCAGCGTGGCCCATTCACCGCCAGCACTTGGTCACCATTGTGATCTACACAGCAAGAAGCA
GCCTAACGACCTGTCTGTTGAAACAAACAAGTTTTCTTTTAAGTGATTTCTTTGTTTCCATTTATAAGGCACCAACTTT
CAAAGGTGTTCTGGGAAACCTTTCTTGTATTTCTTCTAAGCAAAATCAATTCAACAGAGAGTTCACGCTTTGCCTGTGA
TCAATGGGGAAGTACCAGTGTAGCTTTCTTTTTTTTTCCTATAGGGCTGCTCATAGTCCTCCGATAGACTTTACAGCTGT
GGGTACAAGGGTTCCCTCTTCTCCATATCCTCCCCAATATTTATTATCTTTTTTTTCTTCTTCATAATAGCCATTCAACA

Fig. 6. 102

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GGTATGAGGTGATATTGTGGTTTTGATTTGCATTTTCCTATGATTAGTGATGTTGACCACTTTTTCATATATCTGTTGG
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CTGATGGAGAAGAAGAAAATTGTGAGGAAAAGGGTCAATGAGGACATTCTTTGCTGTTTTGCAATTTTCCAATGAGCTG
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GATTGGTAGTGAGTAAAAATATATGTGTGTCTAATGTGTAAGTCTCAAATTATTAACATTTCTCAACAATGTATTTCCA
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CCCCTTTGTGTATAATACTATGAAATGATCTAACTGTTTGGTAGTCCTGGACATGGCTATGTTCTTTCTGCTGTCAGAA
GCCCTGAACTACAGAGTTAAAAGTAATGTTTTTTCTACTGTAATCACATCGCAGCACATTATCTGAACCATTTTAAAAA
TATTTCCCCAGAGTCACTTAATCTTTCTGAGCCTCTGTTCTTTTATTCATAAAATGAGAACAATAATACCTACCTAAAA
GACTGCTTGTGAAGAGTATGATTATGTGTGTACAAAGCATAGTGCGCAGTAGACAATTTTCAAATCTTTCTTTCTTCTC
TCTCCTTTCCTTCCTTCTTTCCTTCCTTCTTCCCTCCCTCCCTCCCTCTCTCCCTCCCATCCGTCCTTCCTTCCATCCT
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Fig. 6. 105!

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TCCTTCTATGTATTGTCCATTATAACTTTGTAATTTCCTTTTGGGCAGGATCCTGTGATTGGTAAATTTTTGTTTTCCA
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TGGGCTCACCGAAGAGTCCTAAAGCAAAGCTTGCTGTGGCTGCAACCTAAGGATTGGAACAAGTCTATAATCCTTCAAG
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AACAATGTGAATGTAGATACCAATATTGCAATTATGACTAATAATTGGAGGCAATTATTTATAGTAGTTATATATCAGT
ATATATATTTACCATTTATACTGATATTAATGACTGTGGCTTTTTAATAGTGGCCACAGAAGTCACATAGCATGGTTTA
GTGAAGTTTGGTTCACATTTGTAGGCACTTCAGAAATATCATCTTTGAGAACACACACACACTTAAGTTCAGTGAGAAC
GTATATAATTCAAAGAATAACATAAGCAATGATTTAAACTTCTATGTATGTTTCAGTCTGCCTCAGTGTTAGCAGCAGG
AAAATTATTCTTTACAATAATGAAAGACATTTCCAGATTCAAAAAAAAGGGAAAAATAAAAACCATGGAAATAATATAT
TTGGAATTACAGGGCTACTAGCAATCTAAGTGTTGTGGAAAATCTTGGTAAATAGTTCAGTGAAATTATATAGATAGAA
GATTGATTGAATAAACCTACTCCAAAGCATTGATATGCCACAGCATTCTTCCTTTGGCTGTGTTCTGCCCAATATTTTA
ACAAGGGGTTGCATCAAAACAGAGTGATGCTGATCAACTCCTGAAAAATATTTAAAGTTAAAAGAAATGCTAAGCAAAA
AGAACAAAGCTAGAGGCATCATGCTACCCAACTTCAAACTACATGCTACAGGAATACAGTAACCAAAACAGCATGTTAC
TGGTACAAGAACAAACACATAGACCAGAGAAAAGAATAGAGAACCCAGAAACAAGACTGCATACCCACAACCATCTGT
TCTTTGACAAACCTGATAAAAACAAGCAATGGGGAAATCATTCCCTATTCAATAAATGGTGCTGGGACAACGGGCTAGC
CATATGCAGAAAACTGAAACTGGACCCCTTCCTTACACCATATACAAAAATAAACTCAAGGTAGATTAAAGAATTCAAT
.ATAAAAACCAAAACTCTAAAAATCCTGGAAGAAAATCTGGGCAATACCATTCAGGACATAGGCACAGGCAAAGATTTCA
TGACAAAAATGACAAAAGCAATTGCAAAAAAAGCAAAATTGACAAATGGGATCTAATTAAACTAAAGAGCTTCTGCA
CAGGAAAAGAAAATATTAACAGAGTAAACAGCCTACAGAATGGCAGAAAATTGTTGCAATCTATCCAGCTGACAAAGGT
CTAATATACAGCATTTATAAGGAACTTAAATAAATTTACAAGAAAAATACAACCCCATTAAAAAGTGGGCAAAATACAT
GAACAGACACTTCTCAAAAGAAGACATTCATGTGGCCAACAAACATATGAAAAAAAAAATCACTGATCTTTAGAGAAAC
CCAAATCAAAACCACAATGAGATACCATCTCACACCAGTCAGAATGGCTGTGATTAAGAAGTCAAAAAACAACAGATGC
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Fig. 6 : 104

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TGCTTATGGAGAAAAACGAATGCTTTTACTCTGTTGGTGAAAGTGTAAATTAGTTCAACTATTGTGGAAGACAGTGTGG
CAATTCCTTAAAGACCTAGAGGCAGAAATATCATTTAACCCAGAAGTCCCATTACTGGGTATATACCCAAAGGAATATA
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ATCAATGACAGATTTGGTAAAGAAAATGTGGTACATATACACCATGGAATTATGCAGCCATAAATAAAGAATGAGATCC
TGTCTTTTGTGGGAACATAGATAGGGCTGGAGGCCACTATCCTTAGCAAATTAATGTAGGAACAGAAAACCAAATACCA
CATGTTCTCACTTATAAGTGATGATCAGAACGCATGGCACATTGGGAGTGGGGAAACAATACACACTGGGGCCTTTCA
GAGGGTAGGAGGGTGAGAGGAGGGAGAGGATCAAGAGAATAGCCAATGGATGCTGGGCTTAATACCTGGGTAATGGGAT
GATCTATGCAGCAAACCACCATGGCACAATTTACCTATGTAACAAACCTGCACATCCCGCACATGTACCCATGAACTTA
AAGTTGTAAAGTAAGAAGAAAAGAACTTGTCTAGATGCAAGTTAATCTGAAAAAAATTTATATTAACTGAGCACACTCC
CAGCCCCAATAACATAGGTGTTTTTGTTTAGCAAGGAATAAGAAAAGAATGGATATTGGGTGGATACCTAATAGTAGTC
TTTGTCAGACTGAATCGAGCAGATATAATAAAAAAGGAATGTGTGACAATGCTATGGTTTAAAAATAACATGATACCCA
GCACTTTGGTAGGCCAAGGAGGGTGGATCACTTGAGGTCAGGAGTTCGAGACTAGCCTGGCCAATATAGTGGAACCTTG
ACTCCACTAAAAATACAAAAATTAGCTGGGCATGGTGGCAGACACCTGTAATCCCAGCTACTTGGGAGGCTGAGGCAGA
AATTACTTGTACCTGGAAGGAAGAGGTTGCAGTGAGCTGAGATCATGCCAGTGCACTCCAGCCTGGACAACAGAGCAAG
ACTTCATCTCAAAAATTAAAAAAAAAAAAGATAGTCCAGAAATAACAAGACACAATCTAACAGGAAAGAAAGGTTAATT
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GTTAATTGCAAGTACTGTATCAAATATGTGATAAGTCTATCTAAAAATCTTCTAAATTGAATCTTTCTTGAGGGAAAGA
AAATCAACATTAATTGAAATATTATTGCTCAGACAAGAAAGGTAGCAGTAGTTCATTGTGCTCAAACAATTGTCAGACC
ACCTATGGAATATCCTGTACAATGAAAAATTACAAGTTTCCCAGAGTGTGTTCTTCAGAGCAGTAGCTCTTAGAACATT
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AGCAGTACTTTTCATGGTCTTTAATATACCAATATGCATTGCAGCTGTCCAAGACAGGAGGCCATTAAGGAAGCCTGAA
GAGAAATGTCCCACCTTAGTGTCCTGAAAGGATATGCACTCCTCTATAGCTATTTACTTTTTACTTTCATCCCCTGACC
TTTCTTCCAGCACTATGGTCCTCTCTTCATCACTTCTCTCTATTTTGATTATTTTCTTCTCCAGTGGCTCCTAACTTCT
TTCTACAAATGCATTTGCTCTGTCTGATAAAAATCATTTTAACTTTGCTTCTCCATCCAGCTACCTTCTTTTTTTACCC
TTCCTGTTACTGCTTCTGGTATTCGATGGTTTTTCCTTTTCCCTTCCCCCATTTTTGAACGATAGACTACTTTTTTTCA
TAGCTCATCTTGCACGTTAGTCTTCCATGAGATATGCCTGTAGGCATGCACAACTAGAGATGTTAAGTAGAGGCAGAAA
GAAAAGAAAGACATCTGGCAATTAGATCTGACTTTATCCATTCTGGCTGTTATAACAAAATAGCATACACTCAGTAGCT
TACGAACAATAGAAATTTATCTCCCAAAGTTATGGAGGCTGGGAAGTCCAAGATCAAGGTGCCAGAAGATTTGATGTCT
GGTGAGGGTGTGCTTTCTGGTTCATAGTTGGCACCTTATAGTTGTGTCTACATGGTTAAAGGGGCGAAGGGTCTCTCTT
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AATATACATTTTGAAGGGACACAAACATTTAGAGCATTCCAAGGTCTTATTTGTTTTTCAGTGGTTAAGAGTTTCGTCA
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CCACCTGGATCATTTAAAATTTTGTTTTTGTGGAGATGGGGTCTCTACAAAAGAGATGTCACTTAGGTAGGTCTCAAAC
TCCTGGCCTCAAGTGATCCTCCTGCCTTGGCCTCCTAAAGTGCTGGGATTACAGGCAGGAGGTACCACACTCAGCCTGT
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CAAACTACAGTGGTTTGGTAACTATATTCATAATTGAAGAAATGGTATTTTTCAGTTACAAGTTAGTAAAAATACAGA
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AGGAGGCAGAGTTTGGAATGAGCCAAGATCACACCACTGCACTCCAGTCTGGACAACAGAACGAGACCCCGTCTAAAAA
AAATTAATAATAAATAAATAAATAAGTTGACGGCAATAGTGGCAGAGTATGAACTCTAACCCATATCTAGGTGTCTCC
AAAGCCTATAATTGGAGAATATTTTGATAATAATGTAGGAGAGAGATTGGTGAGAGAATTAGAGATCACCTTGTTCATC
CTCTTTATTTGATAGATATGAGGACACTGAGAACTCAAAGAAGTTAGGTGACTTACTCCAGGTTACACAGTTTATAGCA
GAGCCAGAAATTGGACTTTGATGCCTTTTTATGTGGAAACATGAGCTTTTATTATTTAGCTCTTCATCTGGTGGAAGTG
GAACACAACCTGAAGAGAGAGGCAATGGACTACACTATGGTTTGGAACAGAGTGTATAGTAATTTCCTATTTCATTTAG
ACAACAGGGATATGCCTGAAAGTGCCTTTACCCATGTCATGCATTTATTCACAATGAACACAAAATTTACTTGAGTAAT
TTTTTTTTTCTCGAGGTATAGTCTCGCTCTATTGCCCAGGCTGGAGTGCAGTGGTGCGATCTCGGCTCACTGCAACCTC
CGCCTCCTGGGTTCAAGCGATTCTCCTGCCTCAGCCTCCCTACTTGAGTAATATTTTAAATGTAACCATAGTGAACTGT

Fig. 6.105

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CCAATACTAAAATTTGTGCCTTTGATAATATTTATATTATGAATAAAAATATGCTCTTTTAACCATGTCCTCATCTATT
TTACCAGAAAGTATCTTGTGTATTGTCAATACAAAGTATCTAATGCATGAATGACTGAGTATGATTGTCTGGTTTTCTT
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GTTCAAGATGTTGACAGAACACATGGTTTTTGTTTTCCTTTACTTTTCCTAATACCCATTAAAATAGTAGAAAAACTAA
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GAAATATGAGTGAGTGGTGTTGATTACTGAAGCAAAACAAAGTCAATTATTGCAGAGAATATTTGTAGAGGGAGCTACA
CCCAAGAAGAAGCCCTAATAACATGGCAGAAGCTAACAAGGCTCCAGTTTCAGATAAGCCAATGCTGATGGACATTGAA
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TGTCATCTTTTACTAATACTCTTTCCCCATGATAGTAGCCCCAAGACAAGTAAGCTGGTCTAGAGGCTCAGTGGATGTT
CAGTGACAGTGCAAGTCATCACCCTTTACTTTTTGCAGATACCACAAAGCTGTTTTTAAGAGTCTCTTGCCATCGTGCC
TATGACTTTGTGCAGTGGAATCAGTTGAGTTCATGGGAAAGCAAATAAAAGGATGGTCTCTGTTTATAGACATAGCCCT
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CAGATACTGAAATTAAACCTGAAAGATGTTGAGAACCCAGAGAAATTCTCAATCATGTTAATAATACTTTCATGCATTT
GACATTTCTTTTTTCCTGCCTCTTCTCTCTACCTGTAGAACTTTTCAAGTTCCATTATTATACCATTGTTGGCCCTTTG
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ATAGAACTTTGTTTTTCTCCTCTGTTAAGGCATGGCTTAACTGTAATTTACTTGAAAACATTATACTGTTTGACAGGAA
AACAGAACCACCTTGAATTATTCTAACCCATTCAGATTTAAGAAAATGCATCGTAGCAAATTAACTGCTATTTGGTTTT
GATCATTCTGTCTGTATCATAGAAATATGGTTCACTAGTGAATCAATGAGAGATAAAAGGTGATCATGGTCTGTGAGAT
AAATGCTGCCCCTCTGAGTTGCTTCCAACGTAGCCAGCTTCAGTGTATCAGTTAGAGTAGTTAATACTAACTGTTGTAA
CAGATAAATACCCAAATCCCAGTTGCTCAAACAATTGAAGTTTATCTCTCATTCATGCTAAGTCAGTTGATGTTTCTGG
TTCACCAGCTGCCTTCCACCTGGTCACCCAGACACCTGGGTTTCTTCCTATGCACCTGTGGCTTCATTTTCCTCTCAAA
CTTCAGACTTCTCTCCCATCAGTATGTGAATAGGGAACCCAGAGATGTTTCCTATGAGCAGGCCCAGAAGTGGCTTATG
TTGCTTCCATTTACCTTACATCACATGTCTATACAAAGAAATGTGGTCTAGCTATGGGCCCAGGGAAGAAGGAATGA
ATTTCATTGACTCTCTGCCACTCGCTCGTTAAGGTCAGATGATACTGCACTCTATCCAAATCAATTTCTTTTTGTAATC
CTTAAAACAAATGTGTTGCATCCATATTATATTTTATAACATATGTCTGTTTCTTTTGGACAGTATTATTCTGTAGACTT
CTTGACTGATACATCCCAAGATCATCTTTTACTGACTTAGGGTTACAGTTTTGGATAAGTTATCCCAATCTCAGGAGTTT
AAATATGCCTTATTGCAGATCTTTTTCCAACTTCTAGGTAAGAAGTAAACTCATTCCTAATCTTACTACTTGGACATAGTC
TCTCCCTGCTACCATGTCCTCACAGGACCTCTTCCTTGGTTCAGTATTGCAACTATCTCCCAGTTTTCCTCTTTCATCT
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Fig. 6.106

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CTGGCTTCTCCTTCTTTATCTCCCTTTTAACTTCTATTTGACCATTAAGTGTTGGCAATGCCTCAGGCGTGGTCCTCCT
TTTTATTGCCTTACACTATACTTCCCATGCCTCTGATTTCAATGATCACTTATATACTAAGGACTCGCCAATTTGTATA
TTCAGCCCATATTTGTCCTCTGGAACTCAGACCTGTATATCCTACCGCTTTTTCAATATCTTCCATTGAGTGTTTCAGA
GGCATTTTAGATCCAAAAATATCCAAAACTGAAATTCTGTTTTCCCCTCACAGCCTGCTTACTGCGACCTGATTTTTCTC
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CCCTCATCCAATCTGTTCTCCATATTGCAGTCAGTCACCTGTTTGAAACAGAAGTCCAACCTTATCACTCACCACACAC
CCTTGCCTTAAAACCAGTTAGCGGTTTCCCATTGTTCTTAAGACTATTAACGTCATTTTTGTGTGATTGTCCAGGGGCT
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GCTTCTTGTACACCGACTTTCCACCCAGTGCCTCAGGGCACTTTCTCATGCTGTTCTCTCTGCCAGGAACACATGCTTC
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CGAGTCAAATGTTGAATATGTATTACAATTTAGGCTGTTCTTGAAGCTTTATTAAATTGAGCACCAAATCTTCTGAAGC
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TCTCTCTTTGTCTCCCCGCTTGGGTCTTTCAAAGACATTTCTGCAGGAGTACAGCTATGAGCTGTTTAGCAGTCAGTAC
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TATAAACAAAACATAGGAGTACATTCAAGAAGAAATGGTGGGAGGGCAAGAACTGAATATCATGAGTTTGGAAAACTC
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TGATATACTTGAAGAGGAGAAAAGAGATGATATCAGGCACATAAATGGAGGATTTGATAAATGTGATGGGGGCTGCAGA
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GTTAAAAATTTAAGGAGAGTTGTGAAATAATCATCTGTAAGCAGGAGTGTGAATGAACTAGGAAGATGTAGTAATATTT
CTGGACAGCCCTGAGAGCTGAAATATTGTAAGAAATAATCTGAGGAAGTTCTGAGTACTTAGACAGTTGAGGTTAGGGG
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TGTACCACCATGGGAATTTGAAAATGCGACCATTTGTACTGGGATATTATCATGTAATATTTTAATTTTACCTCTTAAT
TATCAAAGTGAAAGATTTCCATGCAGTATTATTTGGTTGTGGGTAATTGTTTAATTTATTTGGAATTTTATAGTGCTT
TTGCTTAGCCACATTTTATTTTGTGGTCAAATTCAAATGCTGATCTAATGGCTTCAGAGTAAAATGAGAAAGGTCAAAT
GGGACTGGAAATTAGGGTTTTCTTCCTGTCATTAGGTATATTTATTTATAAAATAACTGGAAAATTCCTCATGTATTTA
ACATCTGTTCCTTCTTTCCCTGGTCACTGCATATCAAGAAGACTTTGACATTGAGGTTTGATACGGTTTGGCTCTGTGT
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ATCATGGGTAAACTTTCCCCCATACTGCTCTCATGGTAGTGAATAAGTGTCACAAGATCTGATGGTTTTATCAGGGGT
TTCTGCTTTTGCATCTTCCTCATTTTCTCTTGCCACTGACGTGTAAGAAGTACCTTTTGCCTCCTGCCATAATTGTGAG
GCCTTCCCAGCCATGTGGAAGTGTAAGTCCAATTAAACCACTTTTTCTTCCCAGTCTCGGGTATGTCTTTATAAGCAAT
GTGAAAATGGACTAATACAGTAAATTGGTACCAATAGAGTGGAGTGTTGATGAAAAGATACCTGAAATGTGGAAGCGAC
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Fig. 6\{16}

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TTTGGAATTGCCCAACAGGCAGAGGTTGAAACAGTTTGGAGAGCTCAGAAGAAGACAGAATAATGTGGGAAAGTATGGA
GCTTCCTAGAGACTTGTTAAATGGCTTTGACCCA...GCCTGCTAGCAATATGGACAATAAGATCCAAGCTGAGGTGCTC
TCAGATGGAGATAAGGAACTTGTTGGGAACTGTAGCAAAGGTGATTCTTATTATGTTTTAGCAAAGAGACTCACAGCAT
TTTGCCATGCCCTAGAAATTTGTGGAACTTTGAACTTGAGAGATGATTTAGGGTATCTGGTGGAAGAAATTTCTAAGCA
GCAAAGTATTCAAGAGGTGACTTGAGTGCTGTTAAAGGCACTCAGTTTTATAAGAGAAGCAGAGCAGAAAAGTTTAAAA
AATTTTGTAGCCTGACAATGTGATAGAAAAGAAAAACCTATTTTCTGAGGAGAAATTGAAGCTGGCTGCAGAAATTTGC
ATAAGTAACGAGAGGCCGAATGTTAAGCCTTCAAGACGATGAGGAAAATGTCTCCAGAGTATCTCAGAGGTCTTCACAG
CAGCCCCTCCCATCACAGGCCTGGAAGCCTAGGAGAAAATGGTTTTGTGGGCCAGGCCCAGGGTCCCCGTGCTGTGTGC
AGTCTAGTGACCAGGTGCCCTGCATCCCAGCCACTCCAGCTGTGACTAAAAGGGGCCAAAGTACAGCTCGTGCTATGGC
TTCAGAGGGCAGCAGCCCTAAGCCTTGGCATCTTCCATGTGGTGTTCAGCCTGCAGATGCACAGAAGTCAAGAATTGAA
GTTCGGGAACCTCCGCCTATATTTCAGAAGATGTATGGAAATGCCTGGATGCCCAGGCAGAAGTTTGCTGCAGGGGCAG
GGCCCTCATGGAAAACCTCTGCTAGGGAAGTGTGGAAGGGAAATGTGGGGTTGGAGCCTCCACACAGAGTCCCTACTGG
GGCACTGCCTAGTGGAGCTATGAGAAGAGGGCCACAGCCTTCAGACCCCAGAATGGTAGATCCAATGACAGCTTGAAGC
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Fig. 6.108

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Fig. 6.509

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Fig. 6.110

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TCTTGAGGAGTATCTTTGTGGCGTTCTCTGTGTTCCTGTATCTGAATGTTGGCCTGCCTTGCTAGATTGGGGAAGTTC
TCCTGGATAATATCCTGCAGAGTGTTTTCCAACTTGGTTCCATTCTCCCCATCACTTTGAGGTACACCAATCCGACATA
GATTTGGTCTTTTCACATAGTCCCATATTTCTTGGAGGCTTTGTTCGTTTCTTTTTATTCTTTTTTCTCTAAACTTCCC
TTCTCGCTTCATTTCATTCATTTCATCTTCCATCACTGATACCCTTTCTTCCAGTTGATCGCATTGGCTCCTGAGGCTT
CTGCATTCTTCACGTAGTTCTCAAGCCTTGGCTTTCAGCTCCATCAGCTCCTTTAAGGACTTCTCTGTATTGGTTATTC
TAGCTATACGTTCGTCTAAATTTTTTTCAAAGTTTTCAACTTCTTTGCCTTTGGTTTGAATTTTCTCCTGAAGCTCGGA
GTAGTTTGATTGTCTGAAGCCTTCTTCTCTCAACTCGTCAAAGTCATTCTTCGTCCAGCTTTGTTCCATTGCTGGTGAG
GAACTGCGTTCCTTTGGAGGAGGAGAGGGGCTCTGCTGTTTAGAGTTTCCAGTTTTTCTGCTCTGATTTTCCCCATCTT
TGTGGTTTTATCTACTTTTGATCTTTGATGATGGTGATGTACAGATGGGTTTTGGGTGTGGATGTCCTTTCTGTTTGTT
AGTTTTCCTTCTAAGAGACAGGACCCTCAGCTGCAGGTCTGTTGGAGTTTGCTAGAGGTCCACTCCAGACCCTGTTTGC
CTGGGTATCAGTAGCGGTGTCTGCAGAACAGCGGATTTTCGTGAACCGCAAATGATGCTGTCTGATGGTTCCTCTGGAA
GTTTTGTCTCAGAGGAGTACTTGGCCGTGTGTGGTGTCAGTCTGACCCTACTGGTGGGTGCCTCCCAGTTAGGCTGCTC
GGGGGTCAGGGGTCAGGGACCCACTTGAGGAGGCACTCTGCCCATTCTCAGATCTCCAGCTGCATGCTGGGAGAACCAC
TGCTCTCTTCAAAGCTGTCAGACAGGGACATTTAAGTCTGCAGAGGTCACTGCTGTCTTTTTGTTTGTCTGTGCCCTGC
CCCCAGAGGTGGAGCCTACAGAGGCAGGCAGGCCTCCTTGAGCTGTGGTGGGCTCCACCCAGTTTGAGCTTCCTGGCTG
CTTTGTTTACGTAAGCAAGCCTGGGCAATGGTGGGCGCCTCTCCCCCAGCCTCGCTGCCACCTTGCCGTTTGATCTCAG
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Fig. 6.⦙ｌｌｊ

ACTGCTGTGCTAGCAATCAGCGAGACTCTGTGGGCGTAGGACCCTCCAAGCCATGTGTGGGATGTAATCTCCTGGTGTG
CCGTTTTTTTAGCCCGTGGGAAAAGCGCAGTATTCGGGTGGGAGTGACCTGATTTTCCAGGTATTGTCTGTCACCCCTT
TCTTTGACTAGGAAAGGGAACTCCCTGACCCCTTGAGCTTCCCAAGTGAGGCAATGCCTCGCCCTGCTTCGGCTCGCAC
ACAGTGTGCTGCACCCACTGTCCTGCGCCCACTGTCTGGCACTCCCTAGTGAGATGAACCTGGTACCTCAGATGGAAAT
GCAGAAGTCACCTGTCTTCTGCGTCGCTCATGCTGGGAGCTGTAGACTGGAGCTGTTCCTATTTGTCCATCTTGTCTCT
TCCTCGATTTTTTTTTCAGGTTTTTAATTTTTTTAATTCATATTTTGATATTCTTCTTAATGCTAGATGTAAGTAAATT
AGATTTTTCCCTGTTAATATTTTATAAACCATTTTTTCTTTTAATTTAGTGAGAGTGACAGATAGAATGATTTACTCTG
AATTACTCTTCCTCTTGAAAGATTAAAGCTTGCTCTTAAGTAAAGTTTGTTATGCCTATCTGTTTTGTCTTCGGTTTAT
TTTTTTCCATGGAGCTGCTTCTTCCTTCTTACAAAATGAAGTTCTGGAGGACCAACAGGTCACTACATGTATTCATAGT
TATATAAATTCTAGCAATGGTAACAGTTTGGCTCATGTTGGGGCCTGGCTACATGTTCTTCATTCCTGTTTAATAAATC
TCATCTTTCTACCATATGCTAGTAATATATGGCTGGAATGCTGGTATGGGAATTACTCCCCTCTTTGCTGAAATAGTTC
ATCTCTTGTGTCCTTTTCCCCTTTTTATTCTTCTATTCTTCTTAGCCTAAGTGATGGTTGTGATTGAATTCAGAAGTTT
GATATTCCTACTCGGTTCATGTCCACCCAAAAGCAGAGAGGAGCCCATCATCATCATTTGTTTTGAATCTGAATCCCAA
GCAGGAAAAATAACTCCAAGGCTCTTACTTAGGCTGTCGTGTCTGTTCTGTCATGCGGTGAGTCAGCAACACTGACTGC
ATTTATTCTTCTGGGAATTGTTCTTCTAGGAGCTGTAGTAGTCATGGTACCTCCCTGCAGGAGTTCATGAATTAATAGG
ACTCATAGTGCAAAGTTTCTCTGAGAATTTTCTATAACACAAACTCCTTAACTTCCTGGTGGGTAATGTTTTCTGGGTG
TTTTTTCTGTTTTCTGTTTTTTTTTTTTTGTCATTTCTTCTTTAGTAAAATGAAAATTGCAAGTAGAAAAGAAACTAAA
AATGGATTTAGTGTGAGGACAGGTTCCTTTTCCTGGCAGGATTGTAGAACACTGGTATTCAGTTGACTGTTTACAATGA
ATATATCTTCTGGTTGGTCATGGCCAGAAGAGAAAATGTCATTGGTTTGTGCCCAAGCAAATTGATTATTAAAATACGT
TGAATATGACCCCATGGTTGCAAACATCCCTTTTCTTAGTAATTCTTAGAGATGAAGAAAGTCTTTTATCTGTTTTCTA
TTTCAACTTTCTGGGTAATCTCTATCCTTTCTTTTTGATAAGTCTCTGTCTTCCCTCTATCAGTGTGAATAATGTTACT
ATTGAAAAACTGGTTCCCTAGCTACCTCAGTCCTGGGGTCTTAAGGTTTCTTCAAAACGTGACAGTCATGTTAAAATGA
AATTCTCTGAATTTATTTAGAGCACTGTTTGTAAAAGTGTAAGTGTTATTATGCAGCCTTGCTTTCCTACCTACCTGAA
TTCTCTCCCTACTCTCCCCAACTCATATTGCCAGCAGTGGGTAGCTGGCAGGATGTTTGAAAGATTTCTTCTTCACAA
TTCAAAAGGCTTGGTCGAGTGACAGGATTCATATCCCAAACCTCAACATCACACAATACTCCCATGTAACAAATCTGCA
CATGTACCCACTGTATTGAAAATAAAATTAAAAAAACTAAAAAAAAGCAAAAAGCAAAAAAACAAAAAACAAAACAAA
ACAAAATCCCTGGTATATATTAAGAGTCCCAGTGACCTACTTCACTTGCAAATTAATGGCCATCTAGAGTGTGGAGTTT
GGTTGAATGATTTATACTGCATTGTGATTCTAAAAATTAATTATTTTTAATTTCTTTGATACTCTAGATCAGTTTTCTA
AGGTAAATGTGAAGTACATCAGTGACTCCAAACTCAATTGAGGTCTCCCGATTTTCTTGGACTGACTATGTATATTTTA
AAGAATTATTTACTTTCATACTTTGCCAAGACCCAGACAGAAAGAAGCTTTCAGTGTTCGTGGCAGCAGAAAAGAATAC
TAATAAGTGTTAGTCATCCACAGTAAATTAAAAATATAATAATAAAGGAAAATTAAACTGCTGTGTGTTTGGGTGTTTG
CTCTATCACCATAAATTCACAACCTCATCTTCTAGTCTCTGAAAAGGAGTGGGTGTGGCAGGAAGCATTTGTGAAATTC
CATTAGCATTTTTAGAGACAATTACTCCACTAAATGATTCAGAATTCAGCTCCTAGCTTCTAAGATTCTACTTAGGCTT
ATGAAGCATTTTATTCTGCAACAGAGTATACATATAATGTAGAATTCCTAGGATTGAAAAACTTGTTAGCTTCCAACCC
AAATTCTCTTACTTTTAGATGAGAAAATGCTGATGATTGGCCTAAAGTCTTATATCTACTGGTGGAGGAGTGGAAATTC
ACTCGTTTACCTTCGCACTGGAGGTAGATTCAAAATACTTCAGAAACGATGTTAGACTAGAGCATAATGCTTGGCCATT
GATAAATAAATAAATGAATATAAGATGTTGTGCCCTGAGATTCAGAATTGGCAGCAAGAATGTGGGTATTATAAAGGTA
TATTTTTCTTCTTGCCTTATTATTGAATAAGAGTACTTGAAAATGGATGAAGGCAACCAGATGTTGAGTCAGATTTCCT
GAGCTTTGAATTTTGGACCTGCATTCATTTCATTTCTCTAGGAAGGTTGCCCCCCTCTTTATTTGTTCAGTGCTTATAC
CAGATTTTATTATACTGGAGATTTCTAAAGCCTCCAACCAAGGAAGATTCCCTGACTTCTGCCTTGCTGCTGCACCACA
CAAATAACCATATCAATTTATATTGCACTATTCTCAACCCCACAAACTCCATTTAGACCTAACAGCTATGGACAGTGGC
CCTAATTTCCTAGGAACATGATCATTTTGTTCTTGTTTGGTGTCTATGCCAAGCTCCAAATCACAAAAAGCTGTTAATT
CCAATTACCTTTTGCAAAATCCATTTTGAAGTCTTCTCACAGATGGATGTATTTAAAACACCTAGAAGGTTAACATCCT
TTGTGAGGCCTCAGCTTTGACTTAGAAAAAGCTCATGCCACAGTGTGTATGTGTGAACCTATTTCCTAAGAGCTGATAT
ATGCCCATGGTGTCTCACTGCAGGGCTGGTGATAAATGATGAAACCATTTAGTGCATAAACTCATATAACCTTTCAGCA
GAATCAATAATTTAGTTGCAATTTAGTTGCATACCTAAAACAATGCAAGGTTGAAGAACCCTGAGAAGAGTGGTAGCTC
CTGATGACAGAAATAGGAGTCTCAGGTATTCTTCTTTTGTTTCCCAAGTGGTATCAAGCATGTGTCTGACTCACTGGTG
CTATTTTACATTGTGAAGGAAGTATGGATGAGGGAAGGCAAAACCAGTGCAAATATTCTAGATCCACTTATTTCCTTCC
CGCTTTCCCTTCCTCTCTTCCCCTCCTTCCTTCTCTTTTATCTCTTCCTTCCTAAATTAATATTTTTGAGGG
TAGGTAAAATAGCTATATTACTTAAAAGTCAAACTTTATAAAAAGGTAAAGCGGAGATGTCTGTCTCCCAATCCCATCT
ATTCCTCCTTCACCCTGTTCCCCATCTATGAGAAATTATATTTATTAGTGTCATGTTTATTTTTCAGTGTTTCTATAT
GTAATACTCATTATATGTTATACAAACACAAATATACATATACATAATTTCCTTCCATTTTTATCAATTTCCTTCCTTT
CTCACTGTCCTGCTCCTAGCTTTAATTTTACCAATTARTAAAACATACTAAAGATTTTTTTCATATCAGTGCATGAAGC
TTTTCCTCTTTTTTCTTTCAACTGCATAATATTCCACTATGTGAAATATCATGGTTTATTTAAATAAATTCTTATCAG
GGAATATTTGAGTTGTTCCAGGGTTTTCTTTCTTTCTTTGCTGTGAAAAACAATGTTACAGTGAATAACTAA
CATCATTTCTCATTTCATATGTGTACAAAGTATCTGTGGGATAGAGTTCCAGAACTGAAATTGCTGCTTGAAAGGGTAA
ATTGTTTCTGTGGATTTGTAGAAATTTCCAGATTTCCCTTTATAGTAGTTACACCATTTTATAATCCTACAGCAAAGTC
TGAAAGTGTATTTTTCACACATCCTTTCAAAAACATTCAAACTTTGGATTTTTGTACATCTAATTGGTGAGAAATGATA
TACTTGTTTTCATTTTCAATCTATTCTTATAGATGAACCTGGACATATTTTCATAAGTTTAATAACCATTTGTTTTTCT
TTCTCGGTAAACCGTATTTCCTTTGCTAATTTTTCACTGGGCATTTGGTGTTGATTCTGTCAAATCTTTACAAGTAG

Fig. 6. 112

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AGAAGTTTGTACTTTGTTTATGATCTGAGTTGCAAATACTTTTATTCCTTCCTTGACATTTATCATTTGCCTTAACTTA
TGGTGAATTTTTACCATGCAAATGTTTTTGATTTTCATGTTGTTCAGTATGTCAATATATTTTATATTCATTACATTCA
TTAGTGGCTAAAACTTCCCAAATGTTGGTAGGAATATGGATGTCTAGATTAGTGAGGCTCAAATGCTCCAGGCAAGATC
AACGCAAAAAATACTTTTCCTAGACACATTAAAAATCAAATTGTCAAAATCAAAGACAAAGTGAGAATTCTGAAAGCAGC
AAGAGAAAAACTACTCATCACATGGAAGGGTACTCACAAAGGCTACAAGCAGATTTCTCAGCAGAAACTTTGCTGGGCA
GGAGGGAGTGGAATGATGTATTCGGAGTACTAAAAGAAAAACAAAACTGCCACATAAGAATAATGAGTCTGTAAAAGCT
GTACTTTAGAAATGAAGAGACATAAAATCATTCCCAGAGAAAAGAAAGCTAAGAGAGTTTATTTCCACTAGACCTGCCT
TAAGAGAAATGCCAAAGGGAGTTCTTCAAGTTGAAATGAAAGGACCCGAAGTAACATCATGAAAACACAGGAAATCCCA
TAGCTAACATCATACTCATGTTGTAAAGCCAAAAGCTTTTCCATTAAGATCAGCAAGAAGACAAGGGAGTTCACTCTCA
CCACTTCTATTCAACATAGTACTGGGAGTCCTAGCCAGGGCCACTAGGCAAGAAATGGAAGTAAAAAATATCCAAATTG
TAAAGGAAGAAGTTAAATTGTCTCTATTTGCATAAGGCATGGTTTTTTATAGGAATCTCTAAAAACTTCATCAAAAAAC
TTAAAATTAATAAATTCAGCAACGTTGCAGGATACAAAATTAACATACAAAATCCAGTTGCATTTCTGTACATTAACAA
CAAACTATTTGAGAAAGAAATTACAAAGACAGTCCGTTTACAATAGCATGAAAATAATAAAATACAGAAGAATAAATAC
AACCAGRGATGTGAAAGATCTGTACACTGAAAACTATAAGGCATTGATGAAAGAAATTTAAAAATACACGAATGAAAAC
ATATTTTATGTCTATGTATTAGAAGAATTAATATTGTTAACATGCCTTTACTGTCCATGTAATATACAGATTCAACCCC
TATCAAAATTCCAATTGCATTTTTATATAAAAATGTGTTGGGTACATTATAATTCTATTCTAGCTATTTGAAATATATA
ATAAATTATTGTTAACTATAGTTTTCCTACTGTCCTATCAATGCTAGAACTTATTCCTTCTATTTAACCATATTTCTG
TATCTATTAACCAACTTCCAGGCTGTGTGTTGACAGTTTTTCTAAGAGTTTTCCCAAGGCAAAGAGTAAGTGGAAGTCA·
ATCTTAAATCACTGTGTCTGGCCGTGACCCGTGATAAAAGATTTAATGGAAAATTTCAGTTTAGATGAGGGTAACAGAC
ATTCATCAGATACTGGGTGCTGGAGATAGATTCAGAGCTAAGTAAGAAATCCTATTTTTGAGATTAAGTAATCTCACTT
CACACCCAGTGACACTGCAACTGTTGATTAAGAGTTTTCCAGTCTCTTTTATCCCCGTTAGCCTACAGGAGCAGGTAA
ATATCTAGTTCCTTCTGGGTCTGAAGATTCTGTAAACTTAAAAGGAGAGATCCTTTCTGATTTGTATATTATTGGATTT
TCTAATGTCCATCACAGCACCGGATTTAACACTCACACACATGCACACACACAGGATATATTGAATGAGTAAATAAA
TAACAGACAGGAGAGTTGTTACTAACTGATCTCTAATATTACCTTATTGTTCTAATAACTGTGGATTTGATATAACTTT
GGGAGCAAGCAAGGTATCAGTCTTAAAGTATAGACAAATTTACTATGTCACCTCAGATACTCAAATAGAGAATATCCTA
TGCCCAAATGGCAAGGCTGAATTGAYCCAAGTGATTAAGTCATACTCTTAAAACTGTATAACCTAATAACCTTTTATTA
AGAATAAGTTCAGAAAAACAGTTCACAATTTCTTTTTACTTAAATTGTATATATTATATGTGCAGTTTTCAGATGTGAA
TGGGATTTGAGATCAGAAAGGCAGGGCTTCATACTGCATTTCTCCAACTTGCTCTCTATTTGTCCTTTAGCAAAAGACA
ATCTTTCCTTCTTTACTTTTTATTTGTTAAACACACACACACACACACACACACACACACACACACACACACACACACA
CACAACTTCATAGCATCCAATGAGCAGTATCTAGGTACACGGCCAATGCAAAGCACTCATTTTCTTATTTCTCCTTCAC
ATGTTTTTCATTCCTATTTTCCATCTTTGCCATGGACAAGACTTCATTCCCTAGCTGTGGCTTTAGAATAACAGTATGA
CTAGATCATATTCATTTTTAGCTTAGCACATTACCTCCCATTTAGTTGAAATGAATATCAACAAGAATTCACACTTCTA
AATGCTTGAAAGCACAATGTCCATCCAAGATCCAAGTGACTGAAATAATAAGAGCCTTGTTATATGAAGAAATACCCAT
TTTCATGATGTTTTCTGAGGCATTAACTGTTACTCTGAAAAGAAAGAAGAGTGTTGGTAAGACTTGGGTATGAGATAAA
ACAATTAGATATTTGCTTATAAACTGGAATGACCTTGTAAATGGCCCAGGAGCAATGTAATTAAATACCATAAAAGCTC
AGAATTCCATTGTGCCCCCTGGAAACATCATAATTGCCTCAGTCATTAGCTTTCAGGGTTTCAGCCTTAAACCAGGATA
TTTCTAGTCTCATCAAAACTATACATTAGCATTCAGTGACTTGTAAGAGTAAACTTCCTTCTAGATAGTCCTGTGTGCT
TATGAAGTACATCCAAATGAACATATGACATGAGGTGGAGTACAACTTAAAGTTGCCTTCTTTCATCCTATATATTTTT
GTTTATTCTTAATGTATCGTGAACTTCAGTAGAGAGCAAACAAGTTGTCTTTATAATTTGAAGTTTTGCTTTCTCTTTT
TGAAAGAAACGCATAATTTGGAGATTCCATCTGTGAAAAAAACTTAATTAAACAATGTTAATTTTTAAAATTCTATTA
TGTATAAGGTGTAAGGAAGGGGTCCAGTTTCAGTTTTCTGCATATGGCTAGCCAGTTTTCCCAACACCATTTATTAAAT
AGGGAATCCTTTCCCTATTGCTTGTTTTGTCAGGTTTGTCAAAGATCAGATGGTTGTAGAGTGTGGAGTTATTTCTGA
GGCTTCTATTCTTTTCCATTGGTCATAAAAACCCTAGAAGAAAAACTAGAAAATACCATTCAGGACACAAGCATGGGCA
AAGGCTTCATAACTAAAACACCAAAAGCAATGGCAACAAAAGCCAAAATTGACAAATGGGATCTAATTAAACTAAAGAG
CTTCTGCACGGCAAAAGAAAGTATCATCAGAGTGAACAGGCAGCCTACAGAATGGGAGAAAATTTTTGGAATCTATCCA
TCTGACAAAGGGCTAATATCCAGAATCTATAAAGAACTTAAACAAATTTACAAGAAGAAAAACAAAGAACCCCATTAAA
AAGAGGGTGAAGGACATAAACAGACACTTCTCAAAAGAAGGCATTTATGTGGCCAAAAAATATATAAAAAAAAGCTCAT
CATCACTGGCCATCAGAGAAATGCAAATCAAAACCACAATGAGATACCATCTCACGCCAGTTAGAATGGCAGTCATTAA
AAAGTCAGGAAACAACTGATGCTGGAGAGGATGTGGAGAAATAGGAATGCTTTTACACTGTTGGTGGGAGTGTAAATCA
GTTCAACCATTGAACTTGTAGATAGACAACTTGTAGATAGACAGATTCCATTGTGGAAGACAGTGTGGTGATTCCTCAA
AGATCTCAAACCAGAAATACCATTTGACCCAGCAATCTCATTACTGGGTATATATCCAAAGGATTATAAACAATTCTCC
TATAAAGACACATGCATATGTCTGTTTACTGCGGCACTTTTCACAATAGCAAAGACTTGGAACCAACCCAAAAGTCCAT
CAATGATAGACTGGATAACAAAAATGTGGCACATATACACCATAGAATACTATGCATCCATAAAAAAGGATGAGCTCTT
GTCCTTTGCAGGGACATCGATGAAGCTAGAAACCATCATTCTCAGCAAACTAACACAAGAACAGAAAACCAAACACCAC
ATGTTCTCACTCATAGGTGGGAGCTGAACAAAGAGAACACAGGGACATGGGGAGTGGAACATCACACACTGGGGCCTGT
TAGAGGGTGAGGGGGCTGGAGGAAGGATAACATTAGGAGAAATACCTAACGTAGGTGATGTGTTGATAGGTGCAGCAAA
CCACCATGGCACGTGTATACCTATGTAACAAACCTGCACGTTCTGCACATGTATCCCAGAACTTAAAGTATAATAAGTA
AGTAAATAAATAAGTAAAAGAAATATGTTAAAAAATTATATTATGTAGATATTTTTGTTATCTTGCTTCATGCCAAGT
GCTTTCATAAATGCTAGACTTCTATTTGTACATTTTGCTTTCATGAGGGTGAAGACAAATAAAGAGTGATCATTCAAAA
ATATTAATGTACGGAAAAGTAAAGTAACTGTAGGGCTTCCATAACAAAAGTACAACAAATTGAATGGCTTAAACAAGAG
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Fig. 6.113

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AAATGTATGTATTGTCTCACAGTTCTGGAGGCTAGGAGTCTCAAATCAAGGTGCCAGTAGCGTTGATTCCTTTTGAGTA
AGCTGAGAGAAAGATCTGTCCCTGGCCTTTCTTCTTGGCCTGTAGATAGACAACTTCCTTCCCTGTATGTCTCTTCACA
TTGTCTTCCATCTGTGCACCTCTCTGTGTTCAAATACCCTCCAACTTAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAGCA
TGTCACTCACATTGGGTTAGGGCCTACCCTAATGATTTCATCTTAACTTGATTACCTCTGTAAAGACTTTATCTCCAAA
CCAAATCACATTCTAAAGTACTGGGGGTGGGTGAGGATTTCAATATATGAATTCTGGGGAGACCCAAAGCAATCCATAA
AAGTAGCCATTTAAATAGAAGTATTTTTAGTCTTTTAACTTAAAGTCTCTCTACARTGTTAAACCATGCCATCTTAATG
TGTGTTATTTAGAATAATCTAAAAAATTATATATTTTCATTAACATTTAGGTTACCTTCAACTTTGTTTTTGATGTTTT
GTAAGCTGCCCATAAGTGACTTTAATCACTCACTGTATACCACTGACAGATCATTTCTCATGAATTCAGTGATGTGCAC
TTTGAGACTCAGGAAAGAGGGAAACTTGGCCTTACTAAAATCATATAGTTAGTATGTGGCAGAGCTGATATTCAACTCT
ACATCTTGCTGCCTCCAAAGTTAGTGATATATGCATGGTCAATATTATTTTAAAATACTCATTTAGTGCTTGCCTTTTT
AATAAAATGCTGATTTTCCTGATTATAAGTGTAACAGGTGATCACTGAAGAAAGTTTAGGAAAAAACTCAAAACTTATT
CAGATTAATATTAAAATTACTCATAATTTCACCATCCATAGATAATTGTTGGTATATTATCTTACAGAAATCGTTTATA
AACATTTTTATTATTCAATTTATATATTTATATCTGCAGAACTACATATGTAGTTTTACTTACCACCAAAACTTGTTGC
AAAAAAACATTTTTACATTGCCCAGTTAAAAATCACCATTAATCTTAGTGGAGAATTATTTTTTAATTATTTAAAATA
AGAATATTTTCATTTTTGCATTTTTCTAGAGAGAGCATGGCATAAGCCTTGTAAAGTTACTCCTGAAATTTAATTTTCT
TTTCTTATTTCCTACTCTCTAGTCCTAAGAGATGAGTGGTTCAAGACAGAGTGGGGGTAAGGCAACAGTCAGCACTTTG
TCTTTGCTCCACTCTCTGGTTCATCTCTGAAGCCTGATTTCTCTACCCAGGCATGAAGATACTTGAATAAGTAGCCTTT
TCCATCCTTTTGCCCTGATATGAAAACTGGTATATCTCTTGAGAGAAGAGCCTCAAAAAATAATCTGAGATAGGAACAA
GTGGAAAGCAGAGATAGTCTCTTCTTTTTAATTCACCCCTCCTTTGGTCACCTCCTGAATATAAAAAGAAGGACCCAGA
ACAACATGGTGTCCACGGCAACCATATCTAGGTATTGCAGAATTTTAGAGAACCCATGAAATCCTTTCAAAGGATAAGC
AGGGGATCTAAATCGTGTCAGATGTATAGCACCCAGCACAGAGCTGAGGGCAGAATATGTGCTTCATGCATCTTTTTTG
TTTGATGGCAATATCACTTCAAGGCTTCTTCAGAAAGATTCCATACCTACCTCTCCTTCTCACTTCTTTATAATCTGTA
ATTTACACACTTACCTCACAACTTTTCTGGGACACATTCAATCTATCAGATATGTAACCACTTGTAAGCTGTTGGTTTG
ATTCTGCAAGTTTTGGTGTGGTAAGACTGGAATCGTACAAATGATATGCTTGGATCAGGTAGTCCCTCGTGGTGAGTCA
TGTCTAGGCTCACCCCGATGCAGTCCAGTCATGCTTCCCCTATGACTTAGTTGTTTGTATTGTCCCCAGAGTCAAGAGT
GTCCCAAGTCTATGAATCTTGCTCCAGAGCATTGTAATTTAGCCTACGGCCTTGTTCACACAAATAAATCTGGCATTAG
TTCCAGCTGGCAGTGATATGGGCTTGGTGATTCACTTTTAGGCCTCATTGGAGTAGAGTTAAAGAAACTTCATTGTCAG
AAAGAGTGAGATCTCTACCATGTGTCTTCTAAGAGTGTTCAGAGTGTTTTTTTTTTTTAGCGTGCAGAAGTCCTAACTT
TATTCCTAGTACCCTCCATTCTCTCTTTTTTTATTGAAACATAAGAATTGTGCATATTTACAGGGTACATGTGATATT
TTTATACATGTATACAATGTGTAATAATCAAATCAGAGTAATTAGGTTATCAGTTACCTCAAACATTTATCCTTTGTTT
GTGCTGGGAACATTCCAAATCTCTTCTTCTAGCTATTTTGCAATATAAAATAAATTACTATTGCCACCTTATGATACTA
TCAAACACTATAACTTATTCCTTCTATCTAACTGTATTTTTTTACACATTAACCAAATTCTCTTCATCTCCTCCCTGCT
TCCCTTCCCAATGTCTGGTAACCATCATTCTACTTTCCACCTTCATGAGATCCACTTTTTTAGCTTCCTTATATGAGTA
AGAACATGCAATATTTGTCTTTTTGTGCCATCGCTTAACGTAATATACTCTAGTTCCATCTATGTTGTTGCAAATAACA
GAATTTCATTGCTTTTTTATGGCTGAATAATACTATATTGTGTATATATTTCACATTTTCTTTTTTTTTTTTTTTTTTT
TTTTTTTTGAGACAGAGTTTCGCTCTGTCGCCCAGGCTGGAGTGCAGTGGCGCGATCTCGACTCACTGCAAGCTCCACC
TCCCGGGTTCACGCCATTCTCCTGCCTCAGCCTCCCGAGTAGCTGGGACTACAGGCGCGCGCCACCATGCCCGGCTAAT
TTTTGTATTTTTAGTAGAGACGGGGTTTCACCGTGTCAGCCAGGATGGTCTCGATCTCCTGACCTCGTGATCCGCCTGT
CTCGGCCTCCCAAAGTGCTGGGATTACAGGCGTGAGCCACCGCGCCCGGCCCACATTTTCTTTATCCATTCATCTACTG
ATAGGTATATAGGTTGCTTCCATATCTTGGCTATTGTGGATAGTGCTGTAATAAACATGGGAGTGCAGATATCTTTTTA
ATACAGTGATTTCATTTTTTAAAATGTATTCCCAGCAGTGAGATTGCTGGGTCATCTGTAGATCTATTTTTAATTTTTT
GAGGAACCTCCATAGTGTTTTTCACAGGAGCTGTACTAATTTACATTCTCACTAACAGTGTATTAGCATTTCTCTTTCT
CTACATCCTTGAAAGCATTGTTTATTTTTTGTCTTTTTGATAATTACCATTTAAACTGGGGTGGGATGATATATCATTG
TGTTTTTCATTTGCATTTCCCTGATAATTAAGTGATGTTGAGGTTTTTTTAAATACACTCATTAGCCATTTGTATGCTT
TCCTTTGAGAAATGTCTATTCATATCCTTCTCTGCTTTTCAATGGGATTATGTACTTTTTTACCTTGTATATTCTGGAT
GTTAGTCCCTTGTCAAATGAATAGTTTGYGAATATTTTCTCTCATTTAACAGGTTGTCTCTTCACTCTGTTGATCATTT
ACATTGTTGTGCTGAAGCTTTTTAGCTTGATGTAATTCCATTGTCTAGTTTTGCTTTTGTTGCTTGTGTTTTTGAGGTC
TTACCTAAAAAATCTTTGCCCAGACCAGTGTTCTTCCAATTCATGAGCATGGGATGGATATCTTTTAATCTTTTGTGTC
CTⁱⁱTTAAATTATTTTCAACAGTGTTTTATAGTTTTCATTCCAGAGGTCATTCACTTGTTGGTTAAATTTATTCCCAGA
TGⁱⁱⁱ TTTTGTGTGTGTGTAAGTATTGTAAATGGGATTACTTTCTTGATTTCTTTTTCAGATTGTTCACTRTTGGTA
TⁱⁱⁱAⁱAAATGTTACTGCTTTTTGTATGTTTATTTTGTATATTGCAACTTTAATGAATTTGTTTTTCAGTTCTAACAGGT
TTⁱⁱGGTYGAGTCTTTAGGTTTTCCTAAATATAGGATCATGTCATCTGTGAACAATGATAGTTTGACTTTTTCTTTTCCA
ATTTAGATGCACTTTGTATCGTTCTCTTGCCTAACCGCTGTGATAGGACTTCCAGTGCTATCTTGAATAACAGTGGTAA
AAGTGGGCATCATTGTCTTGTTCAGATCTTGGAAGAAGAATTTCCATTTTTCCCCATTCAGAATGATGCTAGCTGTGG
ATTTGTCATGTAAGATCTTTATTGTTTCGAGGTATATTTCTTATGTATCCAGTTTGTTGACAGTTATTATCATGAAGGA
ATGⁱⁱⁱAAATTTTATCAAATGCTTTTTCAGCATCTATTGAAATGAGCATATGGTTTTTGTCCTTCATTCTGTTGATACAA
TGTRⁱⁱⁱACATTGATTGATTTACATATGTTGAATTATCCCTGCAACCCTGGGATAAATCCCACTTGGTCATAATGAATAT
TTTⁱⁱⁱATGTTTTGTTGAATTCAGTTTGCTTGTATTTTGATGAGGATTTCTGCATCTGTGTTCATCACTGATACTGACCT
GTAGTTTTCCTATTTTGTTGTGTTTTTATCTAGTTTTGGCATCAGGATAATCCTGGCCTTTTAGGATGAGTTTAGAAGT
ATAGTTTAGAAGTATTCCCTTCTCTTCAATTTTTTTGAAACAGTTTAAAGAGAATTGGTATAGGTTATTCTTTAATCGT
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Fig. 6.114

120/375

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TTGGCAGAATTCAGCAGTGAAGCTGTCAGATCCTGGGCTTTCCTTTAATAGGAGGCTTTATTACTACTGCAGTCTCATA
CTCATTACTGGTTTTCAGGTTTTCTCTTTTTTCATGGGTTCAATCTTCATGGTTTGCATGTGT AGAAATTYATCCAT
TTCTTTTAGGTTTTCCTATTTGTTGGTGTATCAATTTTCATAATCTTGTCTGTTGCTTTTTTTTTTGTATTTCTGTAGT
ATCAGTTGTAATATATTCTTTTTCATCTCTGACTTTATTTTTTAGACTTCTGGTTGGTGTAGCTAAAAGTTTCTTGATT
TTTTTTTCAAAAAACCAACTACATTTTGTTGATCTTTTGTATTTTGTAGTTCAAATTTTATTTATTTATGCTTTGATC
TTTATTATTTCTTTCCTTCTACTAATTTTGAGTTTGATTGTTCTTGCTTTTTTAGTTCCTTGAAGTACAATGTTAGGTT
GTTTGTTTGAAGTCTTTCTACTTTTTGATGTAGGTGTTGATTGCTACAAATTTCCATCCTAGAACTGCTTTTGGTATAT
CCTCTAGGTTTTTGTATGCTGTATTTCCATTTTCATCTGTCCTAAGAAAATTTTTTAAATTTCCTTTTTAATTTATTCA
TTGACCAATTTGTTTTTCAGTAGCATGTTGTTTTCTTTCCATGGATTTGTACAGTTTCCAATGTTCCTTCTCTTATTGC
TTTATAGTTTTATTCAATTGTGGTCAGAAAATATACAGGATATGATTTTGACTTTTTTGACTTTTTAAAACTTGTTTTG
TTACCTAACATATGGTCTATCCTAGAGAATGTTCCATGTGCTGTTGAGAAAAATGTGCATTCTTCAGCTATGGGGTAAA
ATGTTCTGAAAATGTCTGTTAGGTCAATTTGATCTAGAGTAAAGTTTAATTCTGATGTTTCTTTGTTGATTTTCTTGTC
TAGATTACCTATCTGCTGCCAAAAGTGGAGTGTTAAAGTCTCCTACTATTGCCTTCTCTGTTGTTTTAACTGTCCTTG
ACTTAAGGTCTGTTTTATCTGATAATTATAGCTTTACTTCTGCTCTTTTTGTTTGTTTGTTTTCCATTTGCTTCTAATA
TCTTTTTAAATATCTTCACATTCAGTTTGTGAATGTCTTTACAGATGAAGTGAGTTTCTTGTAGACAGCATATAGTTGG
ATCCTTTTAAAAATTCATTCAGCCACTCTGTCTTTTAACTGAAGAATATAATTCATTTATATTCAAGGTTTGAATATTA
TTGATATGTAAGGATTTAGTATTGCTATCTTGTTTACTTGTTTTCTGGTTGTTTTGTAGAATCATTTCTTTCTTTTTCTC
TCTCTCTCTTACTTTCGTTTTTGTCACAAAATAATTTTCTCTATTAGGACATGTTGATTTCTGGCTACTTATTTTTAGT
GTATCTATAATAGGTTTTTGCTTTGTTGTTACTACAAGGCTTATAAAAATATATTTATAACAGGTTATTTTAAACTGAT
AATAGTTTAACTTTGATGCAAATAAACAAAAATAAACTCTACATTGTAATACCACCCCCTCCCACATTTTGACTTATTG
CTGTTTCAATTTACATATTTTTATATTGCCTAACTCTTAATCAATTGTTGAGATTATTTAATAGTTCTTCCTTTTAGCT
TTCATACTTAGGATATAAATTGTTTACTTACCCTAATTATAGTATTATAGAATTATGCATTTTTCTGTTTTTTAATAGT
GAATTTCATACTTTAGGTTTTTTTGTTTTGTTTTTTGTTTTTTGTTTTTTACACATTAGCATCCATTTTTTTTTTCAGAT
TGAAGAATTCCCTGTAGCATTTTTTGTAGGCGAAGTGTCGTGTTGTAAATTTCCTCAGCTTTTGTTTGTCTGAGAAAGT
TTTTTATCTCTTTTTTCCTGTTTAAAGGATAGTTTTCCTGGGTATAGTATTCTTAGCTGGCTTTTTGTATTTGTTTCCTT
CAGCACTGTGAATATAGCATCTCACTTCTTTTGGTCTGCTAAGTTTCTCYGAGAAATCCGCTGAAAGCCATATTAGAGA
TCAGTTGAATGTAATGTTTCTTTTCTCTTGCTGCTTAAAGTATTCTTTCTTTGTTTTTGATTTTTACTAATTTCATTAT
AATATATCTGGGTAGTTTTCTCTTTGAGTTGAATTTGTTTAATATCTTTGAGCTTCCTGCACTTGGATAGTGTTGGTTT
TCTCCAGATTTCAGAAATTTCAGCCGTTACTTTTTGAAATATGCTTTCTAGACCTTTTCCTCTTTTACATAATATCTGT
TGGAAATTTCTATTATGTGGAAGTTAGTTTGCTTGATGGTGTCCCCTTATTCTTCTTTACTCTTTTTTTAAATTCTTTT
TCCTTTTGCTTCTCTGACTGGGTAATTTCATTTATTCTTTTTTTCAAGCTCACTAATTCTTTCCTCTTTTTGATCAAGTC
TGCTATTGAAGCTTTCTATGGAGTTTTTCAGTTCAGTTATTATATTGCTTGCTTCTAAAATTTGTTTGCTCTCTTAAAA
TTGTTTTTATTTTTGTGAAATTTCTTATTTTGTTCCTGGATTGTTTTCCAATTTCAATCTGTGTTTATTGTGATTCTT
TCAACTTCTTTAAGAGAATTATTCTGAATTTTCTGCCTGACATTTTGGAGATGTTGCATTCTTCTGTGTCTATTGTTGG
AGTTTTGTTGGTTTCTTTGAGTGGTGTCATATTTCCCTGAGTTTCCACAATCCTTGTGTCTTTATGTTGATGCTTGTGT
GTTTGAAGAGTCAGCTACCTCTTCCAATTATTGTGGGGGTTTTTTTGGTGGTGTTAGGCCTTTATTCCTTAGTAATGAA
ACTTCAATGTTGGCCTGTTATTTCTTCCCATTCTGGGAGTCTTATAGTGTGCACTGGTACAAAAACACTTTGCTGGAAC
TAACTTGTTGCCCTGCCATTGTTTCCCACTCTGGGGATGTTTTATAGTGAGCACTGGAAGTTAAAAGCTGTCCTGAAAT
TATATTGCTGCCCTACAGTTTTTTTCCCAGTCTGGTAAAGACAGGTGAGTACCAGAACTCAGTCCCAACTTTTAGTTGTT
TCTGGGCCAGGGGAAAGCTCCACCTGGGATTTATAAAAAATTCAGCCACAGATTGGGCCTTTGCTTGAATGGTGCCCC
CCCTTGCAGCACTATGGCATCATCCAGTCTCTTCAACATGGCACCCCCACTGATTGAAGCACAGAGTTGCTGCCTAGAT
CTGCATGCCAGTCTTTGAGATTAATGTCCTGTGCCTTGTCTTCCAATCAGCCCTCTGGGATACCCAATGGTTCCCATAT
GATGGGACTGGAGTGGGCTTCCCATGAAGATTCCCAGACTGATGGGGAGATTGAACATTTCTCCTGCTCTCTTGAGTTG
TCAACCATGTGTTTGCATTTAGGCAGCAGATGGAATTATTGCAAATCATTTACTTTATAACCATTTGTCCCCTTATACT
ATCTTACAATCTACTTTTCATATAATTGTGATGAGTTTTCTGCTGTTATGGAAAATTCCATTTATGGCTAAAGCTTTTA
GTAAATGCTATGAAGCTTGATGACTCTCTCTACTGAATGGGAATAATTATGCATTAGTTCATAAAATGTTCTTATGAGT
TTGTTTAGTCACTAGAAAGGCTAGAATACTGATGCTTAAATATAAAAAATATATAGTTTAAAGAGCTATAATATCTAGC
AGTTCTTTTGTTTCTAAAGCTCCCTGGACATTGAGTAACAAATAAGTTAATAAAGCAATAGTTTGTTATTTGATTTTAT
ATCAGAAAGCCTAAATAGTATTTTAAAAATAAAGCTTCCATATATTCAATCTTATTGGAAAAAGTATTTGAAAAGCTCA
ACCTCAGCTTCCCCTGTTGTTTTATCTAGAATATTTAAGACTTTCTTAACATAAATCATCAAAAATATGTATTTTTTCT
AGTTGAGATTTAACTAGATTAATTCAGTGTGTGGTATTTTTGGCTATCATAATAAAGGTATTGCATTTCCAAACTGCAG
AATATTTGCACTGAGTAAAATAGATGCCATTTTTTTACTAAGTAATTAAAACATTGTCAGAGGATATGCTTTTAAAAAT
TATCACAATAGCTGTAAAGTCAATTGTGTTAATATTGGCAATCTGTCCCTTCTATTTAGAATTTTGTGCCTTAGTCTCT
GATAATTGCATCATTTACAAATCCCCTCTTTACATAAACATGGCCTTTCACCTAACCCTCTAAATTAATCAGCCCTCTT
TTGATTACAAGTGACAGAAACTCAAATCAAAATAGCATAAGCAATTTAAATAATAACAACAATAAATGATGAAAGTTTA
TTAGTTATCTTAGCTGGAAGGATTTTTAAAGATGTAGAAATAAAATGAGTAACTGTAGAGTCAGTGCCTCAGAGACTCA
CCTCGTCATGAGTAGGACTTTTTCCCTCCATCACTTTCTTACCTCTTCTTAGCTACATCTACATGAAGGCATTCTCCAC
TTGGAAAAAAAGACGGAGGCACAACCTCCCATAGAAAAGAGTTTCTTTCTCCCATTATCAATATATCTGCTTGGATTCT
GCTTGGCTATTTGCTTGGATTAAATGTTTACTTTTTAGATACATTACTGCTAATAGAGAATGAGAAAATAAGACTGGCT
ATGCCTGGAGGTTAGGAAGTGAGGAACAAGAGAGGGGCTTCGCAAGAATCATGTGGAATGATTGTGTGATGCTTTCTGA
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Fig. 6.115

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AAGGACATGAGTCTAGGCAGTCAAAAATACCTGGCTAAACACCCTAAGGCTATATTCCCTCTATATATTCCCTATTCCC
TATATTCCCTCCCTTACCCAAACTACCCAGCATCTGCTATTCCCGAACTCTTTTCCCCTCTCATTTTTTTGCATTTCTA
TTGGAAGATTAACCTGTGAATAATCTTTTCAGGTTCCTACCTTTAAAAATACAGGACAAAGACAACTGCTTCATTATAC
TCTTATCTCAGTGGAGTAAAGAAAGTATCTAGAGTTCTGTCTCTTTTTCAAAGTAGATTATTACTCTTAGGAAAAATAA
AATGGGCAAATCAGAGTTTATATTCAAAATATCTCGTAACACCTAAATAAGGCATGGACGCTGAACAATCAGAATTGGT
CTTTGTGTAGACATTGCATCCTGGAGGCCCGTATTATATCAATCAGGCTTAAATTTCCTCATTGCTGGATCATTACATG
GTCCTACTTATCTGGCTGATGGGGCAGGTTAACAGTTGAGGGCTAGGGGAGTTCAAAGTAGTAATTATTTATCCACCAG
GACAGAAACTATGTAATATTTTGGTAATAATACATGACTGCTGACTAACCAAGGCGGGTCACTATGTAGGGGTGCAGGA
TTAACAGTTACTTAATAGAATTAGCCCTGCAGTGACAAACTTTAAGAGAGTTACGTGTTTGGAACATTGATTTAAAATG
TTCTTTTAAGTAGTTGCTTTGTAGCTTGCTTGTTTTGTTTTCTTTTTCCCAAATGTACTTCAAATGGGTCACTGAATAT
CTTCAAATTGTATGTAAATTTTGGAGGGGAAGTGGTATATGCAGATTTTTTTTTTTCTGGAAAGAAGACCAATCACTTT
TGCCAAATTCTCAAACAGGCCCATATTTCTCTCACACACATGCGTGCGCATGCGCACACGCGCGCGCGCACGCACACAC
ACACGTTAAGATGCGAAGAATCACTGTCATAAAATGTCATATTTCTAATTTCTGTAAATGGGGAGAATTCCTCAAGCTT
ATATAAAAGTGATGTAAATAATTCAGTTACAATTATAAAAGAAATGGTATCACAGCAAATCCCTTTGGCACCCCACTGA
AAATTTCTCTCTGATGTTTAAAACACCATTTCTATGCTTACTCTTACAGCTGTATTAATAATCAGATAAGGGGTTTATT
TCTTGTCTATTTTGCTTAGCATATTACCAAATCTGGTTCATATGAACTCATGGGTAAAATCTAAAAATGTTCATGAATT
ATTCTTCATGAATTTTTCACAAAGCAATAATGTATGAAAAAGTGAACAATATAACATCAGCCAGAGACGCCTCAGTTCG
CAATTTACAATGAAGAAAACTGGCATATACTTTAGAGGGAATTTCTAATAAGCCAAAAGGTTAAAAGCTAAATGGGGTC
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GAAAGCAACTGTCAAGGAACAAGCCCATTATTAAGACTGTTACTTAAAGTTCTAAGGGAACTTCCTTAAACCGTTCTAT
GTGTACTTTATTTATCTTGAAACTTTTTGGGAACAGCTGTATAACAGGAAGTAGTATTTCTGAGGCCCAGGAAGTGAGT
AAGGGCTACACAAATTGGGGCTAATGACTCTATTTTGAAAAACAAACAAACAAACAAAAAAACTAAAAAACTTCTCAGG
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TCAGGCTTTCCCATACAAATTTATTTTGCCAAGTTTGCTTTGAAAGAATATTAGATTTTGTAGAGTTTTACATTATTAA
AATCGTGAGTGGCTTCTTTTGCTTTTCACTGTTGAGTGTTTTTGCTACAGCCTAACCTATCTTTTTCTTATTGCTTGTT
TTCTTTGTATTTGCTACTCAATATAGCTTGTCAAATGTGTAATTCATGTCATATACAGTGCTGTCTAATATAATTCATG
GGTATAAAGTGAGAGAGAAATGTAATTCTAAAGTTACTTGAGGTGGACCACTAAAAGGTCCTGAGTTCATTTATTTATT
CAGCAAATATATATTGTCCAATTGCAAGCTAGGCATCAAGTATATTCAAGGTGGTAAATAAAATGAGCCTGATCACTTC
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TGGCAAGAATGAGTAGGAGAGAGAATGGCACAAAATGAAGAACAAGAGGGCAGTGTGGCAGCACTACTGCTGATACCCA
GATATGTGACATTCCACCTCATTCCAGACACAAAGTAGAATTGCACTTCCTGGCTCCCTTGTGTTTGGGTGGGGCCCTG
CTACTAGATCTAGCCAGTGAGTTGTGAGCACAAATGATGCCTGTAACTTTCTGGTCAATTTCTGAAACTAAAGTATGTA
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GACTATAGTGTCCTAGTCAGTTTGGGTTGCTGTAATAAATTACTATAGATTAGGTGGCTTAAGCAACAAACATTTATCT
CAGTTTTGGAGGCTGGGAAATCCAAGATTAAGGTGCTGGCAGATTCAATGTCTAGTGAGGGCACTCCTCTTGGTTTGCA
GGTGGGTGTCTTCTTGTTATATCCTTACATCGTGGATAACAGAGAGAGGAAGCAGGATTTCTTATTTCTTCTTATAAGG
GTATTAATCCCATTTATTAGGACTCTGCCTTCATTACCTAATTACCTCCCAAAGGCCCTACCTCCTAATGTCATCACAC
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GAGGAAATGTCCCTTTCAGAGACTGAATTGGGAAGTAAACCACTTTGTGTAATATTGCAGTAGGTACTGCTATAGTGAG
GAATGGAGAGTGGATGGGGAGAACTCTCCTTTGGTAAGAAGAAGGAAGAAAAGATGTAAAGTAGTGCCTATGATCCTG
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TGTGCATTGAGGAATTGTCCAGAGGGACTAGGGAAGCCCATCAGACTAATGGCAGACCTCTTAGCAGAAACTCCACAAG
CCAGAAGAGATTGAGGACCAATACTCAACATTCTTAAAAGAATTTTCAAGCAGAGAAAACTCAGGATTAAGAAACTCAC
TCAAAATCCACACAATTTCATGGAAATTGAACAACCTGCTCCTGAATGACTCCTGGTTCAATAATGAAATTAAGGCAGA
AATCCAGAAGTTCTTTGAAACCAATGAGAACAAAGAGACAACATACCAGAATCTCTGGGACACAGTTAAAGCAGTATTA
AGAGGGAAATTTATAGCACTAAATGCCCACATAAGAAAGCTGGAAATATCTCAAATCGACACCCTAATATCACAATTAA
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AGGAGATAGAGACATGAAAAACTCTCCAAAAAAATCAACAAATCCAGGAGCTGTTTTTTTTTTTGAAAAAACTAACAA
AATAGATAGACCACTAGCTAGATTAATAAAGAAGAAGAGAGAGAAGAATCAAATAGATGCAATAAAAAATGATAAAAGG
GATATATATCATCACTAATCCCACAGAAATACAAACTACCATCAGAGAATACTATAAACACTTCTATGCAAATACACTA
GAAATCTAGAAGAAATGGATAAACTCCTGAACACATACACCCTACCAAGACAAAACCAGGAAGAAGTCAAATTCCTGA
ATAGGCCAATAGCAGTCTCTAAAATGGAGGCAGTAATTAATGGCCTACCAACCAAAAAAATCCTAGGACCAGACTGATT
CCCAGCTGAATTCTACCAGAAATACAAAGAGGAACTGGTACCATTCCTTCTGAAGCTATTCCAAACAATTGAAAAGGAG
GGACTCCTCCCTAACTCATTTTATGAAGCCAGCATCATCCTGATACCAAAACTGGTAAGAGACACAACAAAAAAGGAAA
ACTTCAGGCCAATATCCCTGATGAACATCCATGCGAAAATCCTCATAAAATACTGGCAAACCAAATCCAGCAGTGTATC
AWAAAACTTATTCATCATGATCAAGCCAGCTTCATCCCTGAGTTGCAAGGCTGGTTCAACATATGAAAATCAATACATG
TAATCCATGACATAAACACAACCAAAGACAAGAACCACATGATTATCTCAATAGATGCAGAAAAGGCTTTAATAAAATT
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Fig. 6. 116

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CAACATCCCTTCATGTTAAAAACTCACAATAAACTAGGTATTGATGGAACATTTCTCAAAATAATGAGAGTTAATTATG
ATAAACCCACAGCCAATGTCATATTGAATGGGCAAAACCTGGAAGCATTCCATTTGAAAACTGGTACAAGACAAGGATG
CCCTCTCTCACCACTCCTATTCTACATTGTATTGGAAGTTCTGACCAGGGCAATCAGGCAAGAGAAAGAAAGAAAGGGT
GATCAAATAGGAAGAGAGAAAGTCAAATTGTCTCTGTTTGTAGACAACATGATTTTATATTTAGCAAACACCATCATCT
CAGCCCCAAAATTTCTTAACTGATAAGCAGTTTCAGCAAAGTCTCAAGATACAGAATCAATGTACAAAAATCACAAGCA
TTCTTTTACAGCAATAATAGGAAAGCAAAGCAGAAAGCCAAATCATGAATGAACTCCCATTCATAATCACTACAARGAG
AGTAAAATACCTAGGAATAGAGCTAACAAGGGATGTGAAGGACCTCTTCAAGGAGAACTATAAGCCACTGCTCAAGGAA
ATAAGAGAGGACACAAACAAATGGAAAAACATTTCATCCTCATGGATAGGAAGAATTGATGTCATGAAAATGACCATAC
TGCCCAAAGTAATTTATAAATTCAGAGCTATGCCCATCAAACTATCATTGACATTCTTCACAGAATTAGAAAAGACTAT
TTTAAGTTTCATATGGATTCAAATAAGACTTTGGATGGCCAAAACAATCCCAAGCAAAAAAAGCAAAGCTGGAGGCAGC
AGGCTACCTGACTTCAAACTATACTGCAAGGCTACAGTAACCAAAACAGCATGGTGCTGGTACTGAAACAGACAAATAG
ACCAATGGAGCAGAACAGAGATCTCAGAAATAACACTACACATCTACAACCATTTGATCTTTGACAAACCTGACAAAAA
CAAGCAGTGGAGAAAGATCTCTTATTCAGTAAATGGTGGTGGGAGAACTGGCTAGCCAAATGCAGAAAACAGAAACTG
GACCCCTTCTTTATACCTTATACAAAATTTAACTCAAGATGGATTAAAGACTTAAATGTAAAACCCAAAACAATAAAAA
CAGTAGAAGAAAACCTAGGCAATACTATTCAGGACATAGGCCTGGGCAAAGACTTCATGACAAAAACACCAAAGGCAAT
TGCAACAAAAGCCAAAATTGACAAATAGGATCTAATTAAACTAAAGAGCTTCTGCACAGCAAAAGAAGCTGTCATCAGA
GTGAACCTACAGAATGGGAGAAAATTTTTCCTATCTGCCTATCTGACAAAGGTCTAATGCCCAGAATTTACAAGTAACT
TAAACATATTTATAAGAAAAAAACAACTCCATCAAAAAGTAGGCAAAGGATATGAACAGACACTTCTCAAAAGAAGATA
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GAATGCTTTTACATTGTTGGTGGAAATGTAAATTAGTTCAACCATTGTGGAAGACAGTATGGCAATTCCTCAAGGATCT
AGAACCAGAAATACTATTTGACCCAGCAATCCCATTACTTAGTACATACCCAAGGGAATATAAATAATTCTACTACAAA
GATACATGCACTTGTATGTTTATTGCAGCACTATTTACAATAGCAAAGACATGGAACCAAACCAAATGCCCATCAATGA
TAGACTGGTTAAAGAAAATGTAGTACATATACACCATAGAATACTATGCAGCCATTAAAAGGAATGAGACCATATCCTT
TGCAGGAACATGGATGAAGCTGGAAGCCATCATCCTCAGAAACTAACACTGGAACAGAAAACTGAACACCACTTATTCT
CACCCATAAGTGGGGGTTGAACATTGAGAACACATGGACACAGAGAGGGAACAACACACACCAGGGCCTGTTCTGGGGA
GGGGATTGAGTGGAGGGAACTTAGAGGATGGTCAATAGGTGCAGAAAACCACCATAGCACACATATACCTATGTAACAA
ACCTGTATGTTCTGAGCATATATCCCGTTTTTTTTACAAGAAATAAAAATAAAGAAAAAAAGAAAATCTCTTGCAACC
AATGAAAATGAAAACAAAATATACCAAAACATATGGGATACAACTAAAGCATTGCTAAGAGGGAATTTCATAGTGATAA
ATACCTATATTAGAAAAGAAGAGGATTTTAATTTAGCAGACTCACCTAGAGAGCACAACTCTGATTTTGTTGAGAAA
GCCTATCTCCACAGAACCTAAAACACACAAACTTTCAAGGTAGGGGGAGATTGGAATCCAAGAAGAATGCAAAAACATTT
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GAACATCTCCCACCAAAGGACTGGGACATCAGAAAGACTGGCACACTCCTAGCAGATCTTCACAGGGAAGGCACTGAGG
GCAGATAGAGGGAAGACACAGATGCTGGGCTGCAGAGGGAGGAAGCTGGGAACCCTGAGCAATGCTAGTGCACACCAGG
ACCGGTTCCTGGTCCCCAAGAACTCCTGGGGATGGGGTGAATTGAACAGGCCAGGAGCGATCCACTCTCGCATGGATCT
CTGGAATCCTGGCATGGGGAAATCCTTTAACCACCATGGACATTTGAATTGGCAGAGAGAGCTGCTTAGAGAAGTGGAA
GGGACAGAAGTCCAGCCAGTGCAGAGCCCAAAGGGTTTGGAGTGGGAGCACCTATAGTGGAGCATGGCCAGGGACACCC
ATCTCCCTAAGCTAGACTTGTTTCCATAGGAGACTGTAGCCCTAGGGGAACTGTCACCTGAACTCTGCAGGGAGGTCYT
GCCCATGAAATGGCAGTCCACCTTGAGTACCCCCTTGGTCTGCTGGCCTCTCCAGGGGCCCCAGCCTGGCTGCAGGTGC
TTGCAGTGCAGTCCCCAGGTAGCTCGTGGGGGCCTGCATTACAGCTCCTGTTCTAGTGGGTCAGGCCTGACTGGCAGAG
TGCTCCAGCATAGCAGTCCCTGCAGACACCAGCATGCTTGTGGCCTCCCACAACTGCAGCGTCCCCCATGCTACTTGCC
CTTCACATACTCACCAATGGCCACCCCCACATTGCTTTGCCACCATGTATTTGCATGGGCAGACTTTGCCTTCCTTACC
CCACCAGGATATGTGCATGCATGCATCCTACCATACCGCTGCTGCCTGCATGAGTGCACCTTGTTGCCAACACCCCACT
AAACCACCATTGTTGTTAGAGCACTGGGTGGCACAGAGCCCACCAGCCCTGCCTTTGCCAGCATTCTGCTCTGGCACCA
ACACTTCCAGCAATGTGAAACTGGACACAGAGAACAGCAGAGTCACCCCTGCCTTGAGCAGCCACCACTGCCTGCATGA
AAACACACAGAGGGTCCACAGTCTTGTGTCCATCAGCACTTCAACCCCATGCTAATACCACCACCAGCACGAATGCACA
CACAGTCACTGGTGGGGGTCCCCGCCCCCAAGCCATGTCGCCATGTGGCCACCACTGCTGCTGTGAATGCCCACACAGA
GGCCAGAACCCCAGCACCTGCTAGCACACTGCCACATCCAACAAGCATGCACCCTGCTGTGTTGCCACTGTCACTGCTG
CTGGCACATGCAAATGAGAACAGATTCTGCTGCCTCTGCCCTATGAAGCTCTTTGTCTGGCACTACCCATCAGAGTGTT
GTGACCAACAGTCCAGAAGTAACTCAGCCCCTCCAGTGCAGCAGGTTCCTAACCTTGAGGAGCCAGAAAACAAAGTTGG
GAACTGATACCAGTCCCCCAGGGTTAGAGCACACAGTCCACGAGTCCTGAGTTGAGTCTTGGTCCCCTAAAATCTTCCA
GAAATTGAGCCAGTCAACTGAACCCACCTTATACCACCATGAAACACCCAAGGTCATGAAATAGAACAAAAGAAAAAAA
TCCAAAGGACAGTAATTTCAAAGATTGGAGGAACAGGCCAGGCGTGGTGGCTCACACCTGTAATCCCAGCACTTTGGGA
GGCCGAGGTGGGTGGATCACGAGGTCAAGATATCAAGACCACCCTGGCCAACATGGTGAAACCCATCTCTACTAAAAA
TACAGAAAACTAGCCGGGCATGGCAGTGGGCGCCTGTAGTCCCAGCTACTCGGGAGGCTGAGGCAGGAGAATGGCGTGA
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AAAAAAAAAAAATTCGAAAAGATTGGAGGAACATCAGACCACAAAGATGAGAAAAAACCAGTGTAGGAACTCTAAGAA
CTCAAAAAGCCTGAGTGTCTTCTTTCCTCCAAATGATCATACTAGTTCTCCAGTAAGAGTTCTTAACTGGGCTGAGGTG
GCTGATATGACAGAAATAAAAATTCAAAATATATATAGAAATGAAGATCATCAAGATTCAGAAGAATGTTGAAACCAATC
TGAGGAAGCTAGGAATCACAATAAAATAGTACAGAAGCTGACAAACAAAACACTCAGTATAGAAAAGAGTGTAACCAAC
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Fig. 6. 117

123/375

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CTGATAGAGCTGAAAAAACACAGTACAAAAATTTCATAATTTAATTGTAAGTATTAACAGCAGAATAGACCAACCTGAG
GAAAGAATCTTAGAGCTTGAAGACTGGCTTTCTGAAATAAAATAGTCAGACAAGAATAAAGAAAAAATAATAAAAAGGA
ACAAACAAAACCTCCAAGAAATAAGATATTATGTAAAGAGACCAAATCTATGACTCATTGGCATCCCTGAAAGAGATGG
AGAGAATGGAAGAAACTTGGAAAACATATTTCAGAATATCATCCATGAGAACTTCCCCAACCTAGCTAGTGAGGCCAGC
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CTCCAAGGTTGAAATGATAGAAAAAATGTTAAAGGTAGCTATAGAGAAAGGACAGGCCACCTACAAAGGGAAGTCCTTC
TGACACAGCAGACCTCTCAGCGGAAACTCTACCAGCCAGAGAGATTGGGGGACTATATTAAACATTCTTCAAAAAAAGA
TATTCCAACTGAGAATTTGATATCTGCCCAAACTAAGTGTCATAAGCAAAGAAGAAATAAGATCCTTTTCAGACAAACA
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CTGCATAGTAACCACCTAACAACATGATGACAGGATCAAACCCATACAAATTAATACCAACCTTCAAGGTAAATGTCCT
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CCATCTCATATGCGGTGACACCCATAGGCTCAAAATAAAGAGATGCAGAAAAATCTACCAAAAAATGCAAAACAGAAAA
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TGACAGTATTAGATCATTGAGGCAGAAAATTCACAAAGATATTCAGGACTTGAACTCAACACTGGACCAAATGGACCTA
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AATCAAGACTAAGAAAATTGCTCAGAACCATACAGTTACATGGAAACTAAACAACCCACTTTTGAATGCCTTTTAGGTA
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AGAATCAGAGCTGAATTGAAGGAAATTGAGACATGAAAAGCCATTCAAAAGATCAATGTATCCAGGGGCTGTGTTTTTG
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AATAAGATAGTAATTACCACTGACCCCAGAAAAATACAAATAACCATCAGAGACTACTATGATGGTTTGTGTACTATGC
ACACAAACTAGAAAATCTAGAAAAATTGATAAATCCCTGGACATATACACTCTCTCAGACTGAACCAGGAAGAAATTGA
ATCCCTAAGCAGACCAATAACGAACTGTGAAATTGGATCAGTAAATTTATCAGTAAATAGCCTACCAAACAAAAGAAGT
CCAGGACCATTCATATCACAGCCAAATTCTAGTAGATATACAAAGAAGAGCTGGAATCATTCTTATTGAAACTATTCCA
AAAAATTAAGGAGGAGGGACTCCTCCCCAGCACATTCTATGAGGCCAGCATCATCCTGATTTAAAAAAAAAAAAAAACAC
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CTGGAGGCATCACACTACCTGATTTCAAACTATTCTACAGGGATACGGTATCCAAAACAGCATGATTCTGGTATAAAAA
CAGAAACATGACCAGTGAAACAGAATAGAGAGCCCTGAAATAAGGCTGCACACCTACAACCATCTGATCTTAGACAAAG
CAAAGGGGAAAGGACTCCCTATTCAATAAATGATGCTGAGAAAACTGGCTAGCGATATGCAGGAGATTGAAACTGGACC
CATTCAGTACACTATACATAAAAATTAACTCAAGATGGGTTAGAGACTTAAACGTAAAACTCAAAGCTATAAAAGCTCT
GGAAGACAACCTAAGCAATACCATTCCAGACATAGGAACTGGCAAAGATTTTATGATGTAGATACCTAAAGCAATTGCA
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TATACACTGTTGGTGGGAATGTAAATTAGTTAAACTATTGTAGAAAGTAGTGTGGCAATTCCTCAAAGAGCTAAAAACA
GAACTACCTGTCAACCCAGCAGTCCCATTACTGGATGTATACGCAAGGGAATAGAAACTGTTTCATCATAAAGACACAT
GCTCACATATGTTCACTGCAGTACTATTCACAACAGCAAAGACATGGAATCAATCTAAATGCCTATCAATGGTAAACTG
AGTAAATAAATTGTGGCACATATACTCCATGGCATACTATGCAGCCATAAAAAGGAATGAGATCGTGTCCTTTGCGGGA
ACATGGATAGAGCTGGAGGCTGTTATCTTTAGCAAACTAATGCAGGAACAGAAAATGAAATACTGCATTTTCTAACTTA
TAAGTGGGAGCTAAATGATGAGAAGACATGGACACAAGATGGAAACTAGGCTTAGCAACTGGGTGACAAAATAATCTG
TACAACAAACTCCATGACACAAGCTTACTTATATAACAAACTTGCACATGTATCTCTGAACCTAAAATAAATGTCTTT
TTAATATCTGGATCCAAATTGTTTTACTGGCCAATGGAAAAAGATTTTTTGCCCAGATGGCTAAATCTTTTGAATAATA
TTTGTGAAAAGACTTTTAAGATTTTTGAATAATATTTGTGAAAAGACTTTTAATAATATTAGTGAAAAGACTTCCT
TACAGAAGGCAAATTAAGTCTTAATTTAATTTGGCAGCTTTTAATGTGGCAATCTTTGATTCTTTATTCTTTTAGATGG
CTGTGTGCACCAATTAAAGAATGCATCCCATTTGCTAAAGAAATTAAAATTTAAAAGAAAAAAATCTTAAACAATTTAA
TACTACATCTCAAGGAAGTAGAACAATAACAACAGACTAAGGCCAAATTAGGCAGAAAGAGGGAATTAACAAAGATTTG
AGCAGAAATACATGAAACGGAAAATTCAAAAATAATAGAAAAAAATAAGACAACTAGGAATTGGTTTTACTGGTTTTTT
```

Fig. 6.118

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ATAAATATAAACTAAATTGGTAAAACCTTAAGTAAACTAACTAGCAAAAAAAGAGCGATGCCTCAAATAAAAAAAATCAG
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GGATTATACACCATGACCAAATGGGATTTGTACTTGGGATGTTAGAATGGTTCAGTGTGTAAAAATTAACGTGAAATTC
CACATTAACAAAATAAAGCATAAAAACATGTGATCATCTCAAGATACAGAAAAAGGGTTTGACAAAATTTAACATCTTT
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TAAAAAGGCATTTAAATTGGGAAGGAAGAAATAAAATTTTCCCTGTTTGCAGGTAACATGATCCTATATATAGAGAAAA
CTCTGAAGATTCAATTAAGAAACTGTTAGAACTAACAAATGAATTTAGTGAAATTACAGAGTACAAAATCAACATACCA
GAATCAGTTGCATTTCTATACACTAACAACAAACTATCTGAAAGGAAATTAAGAAAACAATGCCCATCTAAACTAGTGC
CAAAAAGAATAAAATAATTTAGTAATACACCTAAACCTAACTAACAAGGTGAAAGATTTGTGTATTAAAAACTACAAAA
CATTGATGAAAGATACTAAGCAAATCAAAAGACATTGAATGTTCATGAACTAGAAGACTTAACATTGTTAAAAATATCCA
TACTACCAAAAGAAATCTACAGATTCAATGCAATCCCTATCCAAATCTCAATGTCATTTTTTACAGAAAGTGAAAAAAA
GCTCTAAAATTTATATGGAGCCATGAGTCAAGAAATAACCAAATCAATCTTGAAAAAAAGAATAAAGCTAGAGACATCA
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TACCATAAACTGGGTAGCTTATAACAACAGAAATATATTTCTCACAGTTCCAGAAGTTTGGAATTCCAAAATTGAGGCA
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CAGATGTCTTTTATTCATGAAGTCTCCACTTTTATGACCTAATCATCTCTGTAAGTCCCCGTCTCCTAATGCCATTCAA
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GACGTTTGATAAAGGTGTCAAGAATACACATAGAGATTGGAGAGTCTCTTCAACAAATGGTACTTGGAAAACTGGATAT
CCATATTAAGGAATGAAACTGGATGCTTGTCTTACATCATATGCAAAAATCAACTTGAAGTGGCTTAAAGATTAAACAT
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ACACCTACTATAGTAGAATGGCTACTGTTAAAAAAAACAGAAAATAGCAAGTGTTGACGAGGATGTACAGAAATTGGAA
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GATGATGATGATGATGATGATGATGATGATAATTTGCTTTAGGAAGGAGCTGCACACTACAAAATGCAACAGAGTTGAT
AATGATAAGATTAGTTGAATTAAGTAATTCTAGACCCCAAATTGATTATTTCATTTCACTTTCTAGGTCAATAGTGTAC
CTTGTTGCAATGAACTGAGTTGGTTGTAAAAAGTCTACTATCTTATCAAAAGATTGTTCAAAAACATCTCCATGTACTA
```

Fig. 6.118

AATACTATCAAGCAAAAATATATTGTGTATAGGGAAATACAGTATTTAAAAAAATTGGTAGAATCTACAATGAACTCAA
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CACTCATAGGTGGGAATTGAACAATGAGAACACATGGACACAGGAAGGGGAACATCACACTCTGGGTACTGTTGTGGGG
TCGGGGGAGCGGGGAGGGATAGCATTAGGAGAGATATACCTAATGCTAAATGATGAGTTAATGGGTACAGCATACCAGCAT
GGCACATGTATACATATGTAACTAACCTGCACATTGTACACATGTACCCTAAAACTTAGAGTATAATAATAATAATGAT
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GCCCTTGAAAAATCAAGCAAATAGATGGCTTCACATTTCATTATCTAACCCCGATATGGCCTGACATTTATACTGGGAG
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GCTTTTGGTGATTCCATCATAAAATCTTTGCCCATGCCTATGTCGTGAATGGGATTGCCTAGGTTTTCTTCCAGGGTTT
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TTTCAGTCTTCTGCATATAGCTAGCCAGTTTTCCCAGCATCATTTATTAAACAGGGAATCCTTTCCCCATTGCTTGTTT
TTGTCAAGTTTGTTGAAGATCAGATGGTCTTATATCTGAGGTCTCTATTCTGTTCCACTGGTATATGTGTCTGTTTTGA
TACCTGTACCATGTTTTGGTTACTGTAGCCTTGCAGTATAGTTTGAAGTCAGGTAGTGTGATGCCTCCAGCTTTATTCT

Fig. 6. 120

TTTTGCTTAGAATTGTTTTGGCTATACGGGCTCTTTTATGGCTCCATATGAATTTTAGAGTAGTTTTTTTCTAATTCTG
TGAAGAAAGTCAATGGTAGTTTGATGAGAATAGCACTTAATCTATAAACTACTTTGGGCTGTATGGCCATTTTCATGAT
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CAAAGGGAATGCTTCTCGCTTTTGGCCATTCAGTATGATATTGGCTGTGGGTTTGTCAGAAGTAGCTCTTATTACTTTG
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CTGGAACTGGGGTAAATCAGATGTCAGTATGGTGGACTGACAATGAAGATGGAGTTGCTTTAGTCTCCACACATGGGAA
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CTTCATTGATAATCCTGATTGATGTAGTTTTACTTTCATCATCTAAATATCTCTAGTTTCTAATTTATTCTGTGTAGAG
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TAATGTAGGATTTAAAAACCAATTCAAAAACAATTGTTTATATATGTGGGAAAATTCCTTTAAGTTTCTGTTGATAGGT
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Fig. 6.(2)

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AAAATAGCTATAGTTCTACCTGGAGGTAGAGAGTCATAATAATATCAAATTAGAGCTCATGTTTGTGTGTGTATGTGTT
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GCAGAGTCTCAGTGAAAACTTGTTTGGGGGTTGGTGCCTCTGATTCTCAGTGCAGCCTGCCAGCCTCATTGGGAAAAGG
TGTTTAATATAGGCAGCGATTTTCTGTATTATGAACTGAAATAGAAAATGCTGCGATGGGTTCATTCAAGGGAGACCCA
ATATCTTACTTACAATGAAGAAATCAGGCAATTTTGGGGAGTAGAAGTGGAAGTGTGACTCATACATATATTGGTATTT
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AGATATATGAAAGTGCTTAGTACAAAGCACATGTGTTGCTATCTTATAGTATGCAGTCACGACTTTCATATATATATTA
CATTTATGGTTATTCCAATATATACATTTTGTGTAGAAGGCTTAGTCTTTCAATTACAATAAGCTATTGGTATAGTTTA
GCCATCTAATGCTAATGGAAAGGAGAATGGGCTCAATTGTTCTAAACAAATTTTATAACTTACTTATCTTGGGCTTTGA
AGAAAGGTATGAGGAAATTAACCACAGAGATGTGTGCACACCTAAGAGACATTGACCCCACTTACTTTCTTTCCTTGCT
CCTTGGCTAAGACTCCTATGTAAGTGCCATTGTTGCTAGAAGTTTGAAGCTTTTGTAAGGTTTTAGTTTTTGTGATTTT
ATATCTTTTTTCAGATATTTAATTAAATTTAATATATATTTGCAATCTATAACACCATATAAATAATCATGATAAATCA
AAAGATATATTGTCTGGTCCCTGGTCTCTCGGAGCTGCCAAGGCTCCTAGCTGCTGGGAAAGCTGATGTATATGCAACT
TGGAACATAAGGCAGTGTTTATTGATGCCATTGATCTTTGTAATGGGCTACATGATTGAACCTTTTTTTTATTTCTTCT
CCTTAGTCTTTGATAATGACTTCACATCTGTAGTTATCTCTTTATATTTGTTGAGTTGCTTTTTGCTGCCTGCAGTATT
CTAGACATTTTATAATTAGAGATAAAGCTTGGAATTTCCTATGCAACATGTTTTGTATTCAGCCTGCTTTCTTCTTCCC
ACAGTTTTGGTCTGGTTTTGTTTATTAGTTTGTTTTCTGCAGGTAGAAATACCTAGGAAAGACAACATCATTTGATAAA
GTATAAAATATGCTTATTTAGGAGAGAATACTTTAAAAGGCTTATGAACTCTTGATGACTATCCTTTAGTTTATAATTA
ACTTATTTTTTTTATTATACTTTAATTTCTGGGATACATGTGTAGAATGTGCAGGTTTGCTACATAGGTATAAACATA
CCTTGGTGGTTTGCCACACCAATCAACCCATCATCTACATTAGGTATTTCTCCTAATGCTATCCCTCCCCTAGTCTCCA
CCCACCGACAGGCCCTGGTGTGTGATGTTCCCCTCCCTGTGTCCATGTGTTCTCATTGTTCAAATCCCACTTATGAGTG
AGAACATGCGGTGTTTGGTTTTCTGTTCCTGTGTTAGCTTGCTGAGAATGATGGTTTCCAGCTTCATCCATGTCCCTGC
AAAGGACATGAACTCATCCTTTTTATGGCTGCATAGTATTCCATGGTATGTATGCCACATTTTCTTAATCCAGTCTATC
GTTGATGGGCGTTTGGGTTGGTTCCAAGTCTTTGCACTGTAAACAGTTCTGCAATAAACATACATGTGCATGTGTCTTT
ACAGTAGAATGATTTATATTTCTTTGGGTATATATCCGGTAATGGGATTGCTGGGTCAAATGGTATATCTGGTTCTAGA
TCCTTGAGGAATTGCCACACTATCTTCCACAATGGTTGAACTAATTTACACTCCCACCAACAGTGTAAAAGCATTCCTT
TTTCTCCACGTCCTCTCCAGCATCTGTTGTTTCCAGACATTTTAATTATCACCATTCTAACTGGCATGAGATGGTATCT
CATTGTGGTTTTGATTTGCATTTGTCTAATGACCAGTGGTAATGAGCTTTGTTTATATGTTTGTTGGCCGCATAAAAGT
CTTCTTTTGAGAAGTGTCTGTTCATATCCTTCGCCCACTTTTTGATGGGTTTCTTTGTTTTTTTCTTATAAATTTGTGT
AAGTTCCTTGTAGATTCTAGATGTTAGACCTTTGTCAGATGGATAGATGGCAAAAATGTTCCCTATTCTGTAGTTTGCC
TGTTCACTCTGATGATAGTTTCTTCAGCTGTGAAGAAGCTCTTTGATTAGATTCCATTTGTCAATTTTGGCTTCTGTGG
CCATTGCTTTTGGTGTTTTAGTCATGAAGTCTTTGGCCAGGCCTATGTCCATAATAGTATTGCCTAGGTTTTCTTCTAG
GGTTTTTATGGTTTTAGGTCTTATGTTTAAGTCTTTAATCCATCTTGAGTTAATTTTTGTATAAGTTGTAAGGAAGGGG
TCCAGTTTCAGTTTCCTGCATATGGTTAGCCAGTTTCCCCAACACCATTTATTAAATAGGGAATCCTTTCCCCATTGCT
TGTTTTTGTCAGGTTTGTCAAAGATCAGATGTTTTTAGATGTGTGGCATTATTTCTGAGGTCTACATTCTGTTCCATTT
GTCTATATATCTGTTTTGGTACCAGTACCATGCTGTTTTGGTTACTGTAGCCTTGCAGTATAGTTTGAAGTCAGGTAGC
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GACAATGGGGTTTTCTAAATATGATCATGTCATCTGCAAACAGAGACAATTTGACTTCCTCTCTTTCTATCTGAATA
TCCTTTATTTCGTTCTCTTGCCTGATTGCCCTGGCCAGAACTTCCAATACTGTGTTGAATAGGAGTGGTGAGAGAGGAC
ATCCTTGTCTTGTGCCTGTTTTCAAAGAGAATGCGTCCAGCTTTTGCCCATTTGGTATGATATTGGCTGTGGGTTTGTC
ATAAATAGCTCTTACTATTTTGAGATATGTTTCATCAATACCTAGTGTATTGAGAGTTTTTAGCATGAAGGGGTGTTGA
ATTTTATCAAAGGCCTTTTCTGCATCTATTGAGATAATCATGAGGTTTTTGTCATTGGTTCTGTTTATGTGATGGGTTA
TGTTTAATGATTTGCATATGTTGAACCAGCCTTGTATCCCAGGGATGAAGCTGACCTGATCATGGTGCGTAAGCTTTTT
GATGTGCTGCTGGATTCTGTTTGCCAGTATTTTATTGAAGATTTTTGCATAGATATTCATCAGGGATATCGGCCTGAAA
TTTTTTCGTTGTGTCTCTGCCAGGCTTTGGCTTCAGGATGATACTGGCTTCATAAAATGAGTTAGGGAGGACTCCCTCT
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Fig. 6.122

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TTTTCTATTGATTGGAGTAGTTTCAGAAGGAATGGTACCAGCTCCTCTTTGTACCTCTGGTAGAATTCAGTTGTGAATC
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GATTCAATTTCTTCCTGATTTAGTCTTGAAAGGGTGTATGTGTCCAGGAATTTATCCATTTCTTCTAGATTTTCTAGTT
TATTTGCATAGAGCTGTTCATAGTATACTCTGATAGTAATTTGTATTTCTGCGGGATCAGTGGTGATATCCCCTTTATC
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TGGTCTTAGTTGTTTCTTGTCTTCTGTTAGCTTTTGAATTTGTTTGCTCTTGCTTCTTTAGTTCTTTTAATTGTGACGT
TAGGGTGTCGATTTTAGATCATTCCTGCTTTCTCCTGTGGGCATTTGGTCCTATAAATTTCCCTGTAAACAGTACTTTA
GCTGTGTCCTAGAGATTCTGGTACATTGTATCTTTGTTCTCACTGGTTTCAAAGAACTTATTTATTTCTACCTTAATTT
CATTATTTACCCAGTAGTCATTCAGGAGCAGGTTATTCAGTTTCCATGTAGTTGTGTGGTTTTGAGTGAGTTTCTTAAT
CCTGAGTTCTAATTTGCTCTGTGGTCTGAGAGACTGTTTGTTATGATTTCTGTTCTTTTGCATTTGCTGAGGAGTGTTT
TACTTCCAATTATGTGGTCAATTTTAGAATCAGTGTGACAAGGTGCTAAGAAGAATGTATATTCTGTTGATTTTGGGTG
GAGAGTCCTGTAGATGCCTATTAAGTCTGCTTGGTCCAGAGCTGAGATCAAGTCCTGAATATCCTTGTTAATTTTCTGT
CTCAGTGATGTGTCTAATATTGACAGTGGGGTGTTAAAGTCTCCCAATATTATTGTGTGGGAGTCTAAAAGTCTCTTTG
TAGTTCTCTACAAACTTGCTTTATGAATCTGAGTACTCCTGTATTGGGTACAAATATATTTAGGATAGTTAGCTCTTCT
TGTTGCGTTGATCCCTTTACCATTATGTAATGCCCTTCTTTGTCTTTTTTTGATCTTTGTTGGTTTAAAGTCTGTTTTA
TCAGAAGTTAGGATGGCAACCTCTGCTTTTTTATTTGCTTTCCATTTGCTTGGTAAATATTCCTCCACCCCTTTGTTTT
GAGCCTATGTGTGTTGTTGCACGTGAGATTGGTCTCCTGAATACAGCACAACAATGGATCTTGCCTCTTTATCCAATTT
GCCAGTCTGTGTCTTTTAATTGGGGCATTTATCCCATTTACATTTAAAGTTAATATTGTTATGTGTGAATTTGATCCTG
TCATTATGATGCTAGCTGGTTATTTTGCCCATTAGTTGATGCAGTTTCTTCATAGTGTTGATGGTCTTTACAGTTTGGT
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GTGGTGACAAAATCTCTCAGGATTGGGTTGTCTGTAAAGGATTTTATTTCTCCTTCACGTTTGAAGCTTAGTTTGGCTG
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CTCCCAGTGAGAGCGTCACAGAAGACGGGTGATTTCTGCATTTCCATCTGAGGTACTGGGTTCATCTCACTAGGGAGTG
CCAGACAGTGGGCGCAGGTCAGTGGGTGCATGCACCATGCGCGAGCCGAAGCAGGGTGAGGCATTGCCTCACTCGGGAA
GCACAAGGGGTCAGGGAGTTCCCTTTCCTAGTCAAAGAAAGGGGTGACAGACAGCACCGGGAAAATTGGGTCACTCCCA
CCTGAATACTGCGCTTTTCTGACGGCCTTAAAAAACGGCACCAGGAGATTATATCCTGCACCTGGCTTGGAGGGTCCTA
CGCCAACAGAGTCTCGCTGATTGCTAGCACAGCAGTCTGAGATCAAACTGCAAGGTGGCAGCGAGGCTGGGGGAGGGGC
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GACATCCACACCAAAAACCCATCTGTACATCACCATCATCAAAGACCAATAGTAGATAAAACCACAAAAATGGGGAAAA
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CAGAGAAGTGCTTAAAGGAGCTGATGGAGCTGAAAAACCAAGGCTCGAGAACTACATGAAGAATGCAGAAGCCTCAGGAG
CTGATGCAATCAACTGGGAAGAAAAGGTATCAGCGATGGAAGATGAAATGAATGAAATGAAGTGAGAAGGGAAGTTTAGA
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CACTAAACATGGAGAGGAACAAATGGTACCAGCCACTGCAAAATCATGCCAAAATGTAAAGACCATCGAGACTAGGAAG
AAACTGCATCGATTAACGAGCAAAATAGCCAGCTAACATCGTAATGACAGGACCAAATTCACACATAACAATATTAACT
TTAAATGTAAATGGACTAAATGCTCCAATTAAAAGACACAGACTGGCAAATTGGATACAGAGTCAAGACCCATCAGTGT
GCTGTAATCAGGAAACCATCTCACGTGCAGAGACACACATAGGCTCAAAATAAAAGGATGGAGGAAGATCTACCAAGC
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GACAAAGAAGGCCATTACATAATGGTAAAGGGATCAATTCAACAAGAAGAGCTAACTATCCTAAATATATATGCACCCA
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AGACTTTAACACCCCACTGTCAACATTAGACAGAGCAACGAGACACAAAGTCAACAAGGATACCCTGGAATTGAACTCA
GCTCTGCACCAAGCAGACCTAATAGACATCTACAGAACTCTCCACCCCAAATCAACAGAATATACATTTTTTTCAGCAC
CACACCACACCTATTCCAAAATTGACCACATACTTGGAAGTAAAGCTCTCCTCAGCAAATGTAAAACAGAAATTATAAC
AAACTATCTCTCAGACCACAGTGCAATCAAACTAGAACTCAGGATTAAGAATCTCATTCAAAACCGCTCAACTACATGG
AAACTGAACAACCTGCTCCTGAATGACTACTGGGTACATAACGAAATGAAGGCAGAAATAAAGATGTTCTTTGAAACCA
ATGAGAACAAAGACACAGCATACCAGAATCTCTGGGACGCATTCAAAGCAGTGTGTAGAGGGAAATTTATAGCACTAAA
TGCCCACAAGAGAAAGCAGGAAAGATCTAAAATGGACACCCTAACATCACAATTAAAAGAACTAGAAAAGCAAGAGCAA
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Fig. 6.123

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ACACATTCAAAAGCTAGCAGAAGGCAAGAAATAACTAAAATCAGAGCAGAACTGAAGGAAATAGTGACACAAAAAACCC
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ATCCCTGGGATGCAAGGCTGGTTCAATATACGCAAATCAATAAATGTAATCCAGCATATAAACAGAACCAAAGACAAAA
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CCTGTTTGCAGACGACATGATTGTATATCTAGAAAAACCCCAATGTCTCAGCCCAAAATCTCCTTAAGCTGATAAGCAAC
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AGCCATGCCATTACTGGGTATATACCCAAAGGACTATAAATCATGCTGCTATAAAGACACATGCACACGTATGTTTATT
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ATGAGAACACATGGACACAGGAAGGGGAATATCACACTCTGGGGACTATTGTGGGGTGGGGGTAGGGGGGAGGGATAGC
AATGGGAGATATAGCTAATGCTAGATGACGAGTTAGTGGGTGCAGCACACCAGCATGGCACATGTATACATATGTAACT
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AGCAAAGAAGAAATGCTTTCTCAGACAAAGGCTGAATCTGTCACCAGTAGACCTGCCTTGAAATAAATGTTGAAAGAAA
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GCTTTCTCTCTGGCTGTCCTTAACATTTTTTCCTCCATTTCGACCTTGGTGAATCTGATGATTATGTGTCTTGGGGTTG
CTCTTCTCGAGGAGTATCTTTGTGGTGTTCTCTGTATTTCCTGAATTTGAATGTTGGCCTGTCTTGCTGGGTTGGGGAT
ATTCTACTGGATAATATCCTGAAGGGTGTTTTCCAACTTGGTTCCATTCTCCCATCACTTTGAGGTACACCAATCAAAC
GTAAGTTTGTTTTTTTCACATAGTCCCATATTTCTTGGAGGCTTTGTTCATTCCCTTTCATTCTTTTTTTCTCTAATCTT
GTCTTCACCCTTTCTTTCATTAAATTGATCTTCAATCTCTGATATCCTTTCTTCTGCTTGATCAATTTGGCTATTGATA
CTTGTGTATGCTTCAGGAAGTTCTTGTGCTGGGTTTTTTCAGCTCCATCAGGTCGTTTATATTCTTCTCTAAACTGATTA
TTCTAGTTAGCAATTCCTCTCACCTTTTTTTCAGTGTTCTTAGCTTCTCTGCATTGGGTTAGAACATGCTTCTTTAGCTC
GGAGCCATTTGTTGTTACCCACATTTTGAAGTCTACTTCTGTCAATTTGTCAAATTCATTCTCCATCCAGTTTTTGTTCC
CTTGCTGGTGAGGAGTTGTGATCCTTTGGAGAAGAGGCATTCTGATTTTTGTAATTTTAAGACTTTTTGCACTGGTTCC
TCCCCATCTTCATGGATTTATCTACCTTTGGTCTTTGATGTTGGTGACCCTTGGATGGGGTTTCTGACTGGACATCCTT
TTTGTTGACGTTGATGCTACTCCTTTCTGTTTGTTAGTTTTCCTTCTAACAGTCAGGCCTCTCTGCTGCAGGTCTGCTG
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Fig. 6 (124)

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GAGTTTGCTGGATGTCCACTCCAGACCCTCTTTGCCTGGGTATCACCAGCAGAGGCTGTAGAACAGCAAAGATTGCTGC
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TCGACCCCTGCTGGGAAGTATCTCCCAGTCAGGAGGCACGGGGGGTGAGGGACCCATTTGAGGAAGCATTGTGTCCCTTA
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CGCCTACAGGCACCTCTTCCCCCAGGTGCTCTGTCCCAGGGAGATGGGAATTTTATCTATAAGCCCCTGACTGGGGCTG
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GAGCCCAGTTTGAACTTCCTGGTGGCTTTGTTTACACTGTGAGGAGAAAACCGCCTACTGAAGCCTCAGTAATAGCAGA
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CTGGTCTGCAGGTTGCAAAGACTGTGGGAAATGTGTAGTATCTGGGCCGGAATGCACCATTCCTCACAGCACAGTCCCT
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ACAAAAGAGTTCCTAGTTGCCTTAAAGAATTAGCTATTGTTTCATCTGTGTGTGGCTTCGTCTCAGAGGGTGATTGAAA
CATGACAAGAAGTGATTTTTTTTTCTCAAGAGAAATGGAGGCTTTGGACATTAGTGTCATAGCAATGTCATGTAGAACA
TACATAGAGAATTCTATGAAGAAATCAGCCAAGCCGTGACTATCGACCAAAATTTTCAACTTTCAACAATGAGATGAAT
ATACTATCTATATATCTAATAATCCAAAGCTTCATTTATGAACATATTGCTTCTCAAAATAAAAATTTGGAGCAGTTTG
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TAGTGAGGAGAGTTTTGTTTTCAACACAGTTACGAAAGAATTCGGGCTCACTTAATTGTAAATAAAAATCTTAAAGAAG
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CACCACCCCACAACAGTCCCCAGTGTGTGATGTTCCCCTTCCTGTGTCCATGTCATCTCATTGTTCAATTCCCACCTAT
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AAATTTGTTTAAGTTCATTGTAGATTCTGGATATTAGCCCTTTGTCAGATGAGTAGGTTGCGACAATTTTCTCCCATTT
TGTAGGTTGCCTGTTCACTCTGATGGTAGTTTCTTTTGCTGTGTATATGGAACTAAAAAAGAGCCTGCATAAGCCAAAA
GAACAAAGCTGGAGGCATCACACTACCTGACTTCAAACTATACTACAAGGCTACAGTAACCAAAACAGCATGGTACTGG
TACCAAAACAGAGATATAGATCAATGGAACAGAACAGAGCCCTCAGAAATAACGCCGCATATCTACAACTATCTGATCT
TTGACGAACCTGAGAAAAAGAAGCAATGGGGAAAGGATTCCCTATTTAATAAATGGTGCTGGGAAAACTGGCTAGCCAT
ATGTAGAAAGCTGAAACTGGATCCCTTCCTTACACCTTATACAAAAATCAATTCAAGATGGATTACAGACTTAAACATT
AGACCTAAAACCGTAAAAACCCTAGAAGAAAACCTAGACATTACCATTCAGGACATAGGCATGGGCAAGGACTTCATGT
CTAAAACACCAAAAGCAATGGCAACCAAAGCCAAAATTGACAAATAGGATCTAATTAAACTAAAGAGCTTCTGCACAGC
AAAAGAAACTACCATCAGAGTGAACAGGCAACCTACAAAATGGGAGAAAATTTTCGCGACCTACTCATCTATATTTTTT
AATGGATTTAACCAGCAAGAAAGACCATGGAGCCACCTAAATTTTTCCAAAATCTGCAAAATGAAGGTGATATAAATAT
```

Fig. 6. 125

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GTCACTTAGACAATATTTCATTTTTGATATAAAATTTTATTTTTATCAGTTTGATAATATGCATAGGACTAAAAAATG
CAGTTGTCTTAAAATATTTTAGGGTTGCTTAGGA  TCACTTTAAAAATAAAAAGTGTCAGAATAAAAGTTGTCTGCTT
ATGTTCCATATTTCAGAACTACCTTTGATTTCTTTAGCTAAACAATATTGCACATAAATAGGTTAGGAATTATAGACTT
TCAGATGTGCTACCTGTCTCCCTTCAAAACACTCATTTAGGTAATGTATTAATTATTTTTAGAATCACCTGTCTTCTCT
GCTTCAGCAAAAACCCCACCAGGGCAGAGATCACTCTCCCTCTTGTTTGCTTTGAATTTTTATGACTTAGCACAGTGAT
TGGCACATAAACATTATTAAAACCAATGAATGAGGAAAGCAACGAATAAGTAAACAAATGGGCCCAGACTAAAAAGTAA
GTTATATGTATTAGTGAGAGTCAATAATAAATTAATGAAATAATCTTTTATTTGAAACTGAAGCAAATTAAGGGAAATT
TATGAACCTTAAATCTTTTTGTAAAGATCTAGTACCAATAAGATAAAAAAAATGCTCATCCTTTTTTCCATTGAAATGT
TATGTTCAAATGAGCTTTGTTTCTATACTTTATACATCAATTAAGCTGAATTCATAGGTGCTAAGCATTTTAACATATT
GTATTGCATGTAATATTCCCTAATGCCCCAAACTTCAAGATTATATAATGGCTTACTCTCTCCCTGTCCCTACCCACCA
GATAGTGTTATCCACATACATTCTCATCTAGTTTTGTTCTGTGATGAAAAACCATATGCGTATCCTATTCCTATGTGAA
TTAACCTGGCATGCAGGTAACTGAAGTGCATGTGATCTGGTCTCTGCTGACATATCACAATGGGCCCCCCTCCTTGCAT
GGAGGTAGCGTTTTTTATAAGACAAAATGTTTTTAAATAGAACACATTTCAGATTTCAGATTTTATATGTATTTTGTGT
TTTTCCTCCCTCTACCCCTTTCCAAAATTATGAATGAAATTCTAGGACCATTTATAGACAAAGCACAGTTTAGTCCGAG
TAAACCAATGGACCCTGAAGTGTCCAAGTCAGAGTTTATTTGCAGAATTTAATTCATTCCCACAGAGATAGGCAACCCA
GGCCTGTGAGGACACGAGGGTAACTAGGCAACAAAGTGCCGTAGTCAGGCTTGTGTTTGCTTTTTGGTAAGAGGACAAC
ATTGACTTCAGTGTCAGGGCATAAAGGAGACTCAGGACTTATTAATTTTTTTCCCCATAATTCTGTGAACTTTGTGAAT
TCCCTAATATTCTTTTAACAAGAGTTCGGAGACATGAGTTTATGTGCCTTCTTGGATATATTCACGGGAGTTTGCAGAG
AAAGTTGAATAAATTATTAGGTTAATGGCCTGTGTAAATTCACCAACACCTTTTCACTATCTCATCAATCATCCTGTTT
GAGTTATTGTGATTCACTCTGCTGATGAGCTCACACCCTTTTTTCCTGATACAGGAATTATTGTACACCAGGGGACTGG
ATTTTAAACAAAACGTATTCCTTAGAATAACTTGAACAATGGATTGGTGGGTCCTTACACTATTATGTGCTGTGTAGCT
GTACAAGTGTGTCTGCATGAGCTTTAGGACATTATTTGAGATATTTTAAGCTATGTGTACCTCATGAACTTGTAGCTGA
TTTTCCTTAGTTCTTTTAAAATATTTTCTCAGAAAACCAACAGTAAAATCTATCAGGTTCACATGAATACACTCATTTG
TGTCATATCAACCCAAAATGAATATGATCTTCCAGGTAATGATGAAGGATGATAACTATAATATTTCCAGCCAAACTTT
ATTTTGAAACATCACTCAGTGTTTCACATGTTTAGTGGCTGTTAAATTCTATATGTCTAAGCAAACTGTGAAGAGCATA
ATTAACTTATTTGGTTGTTGTGTATCTTTTAAAATCACAGTTTGAATCTGCTGGGAATGTTATAGTGGCACTAGTAGCA
AAGGAATGGCAAGGATGTTAAGACTTTCTCATGCTAAGACCCCAGCTTGGTATTGAGTTTTTAGGAGGGGCCCGCATGA
CATCTCTAATTGCTTTGGCAATCCAATTATTGAGACAAGTAAGTGAGCCCATCGGGTGCCTCCCAAGGAATTAACTTGG
TCATTTGATATAATGTTGCACTCTCCCTATTTGGGAGGAGGGGTGGCCACCCTCACCTCTGCCATTGAAGATTAACCCA
CTACAAATTTTCAAAAATATAATATCTAATTTATTACACAGACTATCATATGGGTTTCATAATCCTTAAGTATCTCTAA
ATTTTATTATTATTTTAGATTTTTAAATAAGACTCCACCTAACTTGATGGAAGGAAAAGGGGGTCTAGAAGCATACAGA
GTTTGGCTCTTGGTTCTCAGAGTGACTCGCCATGTGAACATGGGCAAGTTATTTAACTTCTTTGAGCCTTTTGAATTTT
AATCTGTAAAACAGAGACAAAAATGCCTTCCTCAGAGGGTGAAAATAAGGATTAAGTACAATAATGAATGCAAAATACC
TGGCAGAGGACCTACCTTGATGTGCTATTCTCAACATTGATATTTAATCAGTATACACAAGTATGCCCTTTTTTTTTTT
TTTGAGACGGAGTCTCGCTCTGTTGCCCAGGCTGGTGTGCAGTGGCGCCATTTCAACTCACTGCCACCTCCGCCTCCTG
GGTTCAAGCAATTCTCCTGCCTCAGCCTCCCGAGTAGCTGGGACTACAGGCATACGTCACCATGCCCGGCTAATTTTTG
TATTTTTAATAGAGACTGGGTTTCGCCATGTTGGCCAGGCTGGTCTCGAACTCCTGACTTCAGGTGATCCTCCCGCCTC
AGCCTCCCAAAGTGCTGGGATTACAGGTGTGAGCCACCGCACCCTGCTGCATGCACATTTTGCCTGGATGATCTTCCAG
GCTACCATGTGGATCTCTCTCATACAGCAGAGGGAGAGTTGTACAGCAATGAGGGCTTCAGAACTGTGTCTGACCCATG
CTGATTGCTCAGTGCCCCTGAGCACAAGTTTTTAAACATTTTGAATCTCACCCATAATATTTCTCATCTTGCAGAGACA
GAGAGCATTGTGGCTATATAATGCAGGTAAAGTACCTAGTGTAACGAATAACATGCGGAAATGATAAATATCATAAGTT
GTATTACTGCCTGTGGTAGTCTGAATAATGGTCCTCAAAAGATATCCAAGTCTTAATTTCCATAACCTGTGAATATGTA
ACCTTATATGTCAAAAGGAACTTTGCAGATGTGGTTAATTTAAGGATCTTGAGATGGGGGGATTTTCCTTGATGATCCA
GGCAGGCCCTAAATGTAATTATAAAGGGGCCTTTTAAGAGAGAGGCAGGAAGGTCAAAGGCAGAAGAAGGCAATGTGAC
AGCAGAAGTAGAAATTGGTGTGAAGCCGCAAGCCAAGGCATGCAATGCTGAAAGCCTCTAGAAGCTGGAAGAAGAAAGG
GATGAATCCCCCATTGGCATCTCCAGAAGCAATTAATCCCACTGACATCCTGATTTTAGCTTAGCTCTGTAAGACTAAT
TCAGGACTTCTGACTTATTTGTGGTGTTTTAAACCACTCAGTTTGTGTTAATTTGTTGTAAGGGCAATGGGAAATGAAT
GCACTTCCCATCCATAACCCTGTCTCTAGCTATGTTGAAGTCTTTACCATCCCCCAAATCTGACTTTTACTTGCATGCT
TCTTGCCTTTACATATCTGCTTTTCCTTCCTGCTTCTTCCTATCTCCCACATCATTGCCTTTCTCACAACTTCTCTACT
CAGCAAATTCCTGCTTATCCTCTGAGAGGGAGGACAAGATCATCTCAGATATAGTTTGCTAGCTAAGAAAAGAGTCCTG
AGACTGCTTTGAAATCTTGTGGTCCTTTATCCATAGCTCATGTAAATAACTCTGAGCTAGCAGTGTATATTATTTTTAT
TTCATCCTGGAACTATGTCTTAGCAATATTTTTATTGGAGAAAAAACTGAGAGCAAACTCTCTGTTTCCCACATGCCTA
TTGCCAAATTTCTCTGGGTGCTTTACTTTCACTGCACTGATTCTAGAAATTGTATTGACAACTGTGTTCTAAGATGACC
CTTTGATTCATCCAGATGGTGTGAAAATGTTCCTCTTCAAAAACCCCACACTGCTTTTCAACATCCTAATGGGTAGAAT
CTCTAAATCCTTAGGGTGTCAACAAAAGCAAAGGGCATATTTACACTTCAGTTGGGTGAAATTAATTAGCATGGAAGTT
AATATTTACAATTAATGGAGATTACCTGACTTTAATAGTGAGATTGTGTAGCTACTTTGTATCACTCACTCTTTTTTTG
TTGTTGTTTTAGAAATTGTGGCAAAAAATGGATAAAATAAAATTCACCCTTTTAAAACATTTTTTGAATTCCTGGAGTG
TAAAAGTGGTAGGATAACCAAAACATCCTCTTTCTTGTTGAACCATCAACAATTGTCTTTTGAAAAGAGGTTTGAATAT
CTTTTCTTTGAAAATCCTCGTGCAAAACTTCACACCATGATCATTTATGAGGTAGTTATCAGACACTGGAGATGAATTA
CTGGGGATTTTTGTGTTCTGGTTCTCAGTATACAACCAAAGGTAGAATATTGACATTGAAAATAAATGTCACTCGTTTC
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Fig. 6.126

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TCAAGGAAAATTTGTAGTTTACCGAACAACAGCACAAATGAAAGCTGAGATACTTTACCAGTGATATTGTAGGTCTCAG
TGAGTAAAAACTCAATTAAAAGTATATTGGGGTGGCGGGGGCACTGTGTATAGATAGACCTGGACATGATCTCTAATCA
CAAATGCTACTTTTAGAAGGGCCAGATTCTTGATACAGAGACATTTTATTTGGTGGCAATAACCATGGCTTGTCCACAG
AATGATGCCGTATTATTCTCCTGACCTAACTTCAAAGAAATAAAGAGTTTGCAAGAAGAACTGCAGTTCTTCAAAGTAC
GCAATATGGATTTCCAAGATGAATGTAGTTTCTCTCTCTGAGGAATTCTGAACAGTGGTAAAGTTTCACAAGTTTATGC
ATATCTTTTGCTCTACATCCTTCCCTAAAAGAACTTGTGGCAACAAACAAAGGGAAAGAAATAACATTTTTTAATAACT
ATGTGGAAAGCTCTATAGTAAGTGCTTCACATGTTACCTCATTTAATTCTTACAACTACCTTATCTAGTAAATTATCTC
CCATTTGACACTTTAAGGAAGCAGCTAAGAGATGTTTCATAACTTTCCTAGAAAAGGTAGGATTTGAATGCAGGTTTGT
ATTATTCCAAAGCTCACAATGTGCTTTACGCAACATCAAAGTAACATATTGCGGGAATGAGTACCTTTCCCATTTAAAA
CAAATGAGTCCTGGAAACTCTTACCCTGTTTAGTTATGGAATGGCTCAGAAAATAGAAAGTGTTGAGATCATCAAAGAG
AGAAGTTAACAAAGAGCATTGTAATCCAGAAATAAGAACGCAATAGAGAAGTAGAAGTTGTGTGGCTAATTTTACCAAA
CTAAATAGCCTGAATTATTCAGTGTGACTATACACATTGATCAAATTAAATGAGCATACCATAGTCTAAAGGGGACGAG
ATTTATATTCTATCCAAGAAGTCATTAATTATGTTTGTACTATCTTCATCATGGTTATCATTTTTCTTAGACATAGCCT
AATCTATAAGATTTTACTGTATTTCCCTGAATTACTAAATTCTTCTATTTTTGAAGTTTTACTAAGATTTTATTGTATT
TCCCTGAATTACTAAATTCTTCTGTTTTTGAAGTTTTACTAAATACTTCAGAAGCTTTACTAATTAAAAAGGTAATTTA
TAATGTTTATCACTAACCAGTTGATAATAAAGCGCTTCCTTATAGCTTCTTAAGATAATAGCTAGAAAACAAAGCTGAT
TTTAATTATTCTTGTAATTTGCTTCAACTTCACTGACAGTCTGTTGTATATTTTCTGCATATGTAATTACATCAGGTTT
TGACTATTTTCTCTGAGGTAAATAATTCCAATTACTTTAACCTGTCATTGTTTCTCTTTCTCTCTCATTTCTTTCTCTC
TCCCTCCCTCTCTCTTTCTCTTATTATAAGACATATTATTTGTTTAATTTTAATAGCTATTCCAGCAGTTAAATATTCT
TATCATAAATTTCTTGCAATTTTTAAATGTGGAATTCCAGTACAAAACTTTGGTTTAGCAGTAGGTAAAAAGATAGATT
GAAATTTATGTAAACATATTATTCTTCATCAGGCAGGGCTAGCTGACTGACATCTGCTTTGGTTTGGATTCAAATACAG
ATTTAAAGGGAGGCTATACATGGAAATTATCCCCAGGCTCCTCCCCGCTGCTGTTCCCAATTTCTGCAGCAGCCTGCAT
GCTTGTATTTAGGATTCACCAATTTTTTTGGCCCTTGTGTGTGCTTGGCTTTTGTTTTGATTACTCTACAATGGCAGG
ATGTTCTGGGAACTGATGTGATGAAGTAAAGGAAAGCAAATAGCCTTTGTCATGTTTAATTTCCCTTAAAATCTATTTG
CCAGTGAAAGGCCTTTAAAATTTTTCATTATAATTAAATGTTTGTTAAAGCAAAAAGCCTCTTCTTGTATCTATAGACT
AAATCAGGGTTTCTTTAATCATAGGTGGTTAATCCTCACTACTCCTTTACCCAAATACATTTCATAAGATGCTAAAGAT
GTCAAGGAAAATTCATATCTCTGACTAACAGAAAATATTCTTTCACAAATAAAATGTGAAGTGTTCTTAGTTGTCCACT
ATTCTTTTAGGTCCTACATTATTCTATATCTATTTTCAACAAAAGCATAAATTATGGGAACATTGAATTAAAATAACAT
GAAGGTCCATACCATGTCCCTGGATCAGAAGTCTCCAAATCATAAAAATTGCAGTGGTCCATCAACTGATCTGTAGATT
TGATATAATTTAAGTCAAAATACTGACTTGTTTATTTTGGGGGAACTTGCCAATCTGGGTCTAAAACTTACACGGAAAA
GCAAAAATTAATAGGCCACTGTAGATTTTCCTTACGAGGTAGTGTGAGTTTTTAAATTCATTTATTTATTTATTTTTGA
ACATTTAATTTAGGTTTGGGGTATATGTGCAATTTGTTATAAAGGTAAACTCGTGTCACAAAAGCATTGAATTTTATG
CCATGTTATCTATGGTTGAAATATAATAGGGAGCGAGTTTATTTTAAAATCATGTGTCTTTTTAAGATTTGATTTATGC
TGACTTCAGAATGTGACGAGTTTTAGAAATCTGAGTGTAGGAATGAATGTTCTATTGGAAGTATCTAGTTTATTGCTTT
CAGCTCTACTGATCATGGATATTAACCAACTCTTGTTTTTGTTTTTTGTTTTTTGTTTTTTGAGACGGAGTCTCGCTC
TGTCGCCCAGGCCGGACTGCGGACTGCAGTGGCGCAATCTCGGCTCACTGCAAGCTCCGCTTCCCGGGTTCACGCCATT
CTCCTGCCTCAGCCTCCCGAGTAGCTGGGACTACAGGTGCCCGCCACTGCGCCCGGCTAATTTTTTGTATTTTTAGTAG
AGACGGGGTTTCACCTTGTTAGCCAGGATGGTCTCGATCTCCTGACCTCATGATCCACCCGCCTCGGCCTCCCAAAGTG
CTGGGATTACAGGCGTGAGCCACCGCGCCCGGCCTTAACCAACTCTTTAAGCAGTGTTTGGCTCATGGACATTGGGGTT
AAGTCTGTGTAATTAGGGCTTTAGGATATTGGGATTAATTAAACCAGTTGCTGCTTTAATCCAGCATTTTAGGCATCAA
TGGTTGATGATGGGAATGATTCTGTAACTTGCATTTATGGTCTAAGAGCTCACTGGTGTATATACCTGGGTTAATTGGG
GATTTTAGGTAGCCAGAATTGGAAAATAGAATGCACAGTATGAGAAATTCCCCCGTCTCTGAGGATGGGGGATGCTGGG
GCCCCCGTACCATCAAGTAGCACATCCACAACTCACTCCATTTACCACATCATCACCATCACCACCACCAACAAGCACT
TGACCTCCATCCTAACTTCAACAATCACCTTCAGTCTGCCTAACTGATGCTATGGAAGAACAACAGAGACTCAGAAGGG
TAAGAGGGTGGTGGTGGCGGGTGGTAGATGATGAAAGATTACTTACTGGTGTGAACCTTCAAGATTTGAGACAAGTCTT
AGTTAATTTGGAAAGTTTATTTTGCCAAGATGGCATTGAGGATGTGCACCCCTAACAGCCTCAGGAAGTCCTGACAATA
TATACCCAAGGTGGTTAGGGCATAGCTTAGTTTTATACATTTTAGAGAGACATGAGACATCAATCAATGTATGTAAGAA
GTGCACTGATTCAGTCTGGAAAGGCGGGACAACTTGAAGCAAAGGCAGGAAGACTGGAAGCGGGAACTTACAGGTCACA
GATAAGTGAGATGAATGGTTGCATTATTTTGAGTTTCTGATTAGCCTTTTTAAAGGAGGCAATCAGATATGCATCTATC
TCAGTGAGCAGAGGGGTGACTTTGAATAGAATGGGAGGCAGGTTGGCCCTAAACAGTTCCCAGCTTGACTTTTCCTTTT
AGCTTAGTGATTTGGGGGCCCCAAGATTTATTTTCCTTTCACACTGGGTATAAGGTACATGATTTGAGTGATGGATACC
CTAAAAGCCCTGACTTCACTGCTATGCAATCTATGCATGTAACAAAATTATACTTGTACCCCATACACATTTATACAAGTA
AAAATAAATAAATAACAAAAACAAAACAAAAATAACAATGGCCCTAGCCATAAAGCAATTTTTAATTTAGCGGGGAGGA
CAGATATGTATCTAACCAGTTATACTGGAAAGTGGAGTGAGCTCAATATGCTAAAATAGAATTATTGTGCATCATAAAT
GTTTAAGAGGAGGGAAAGGGTAATCTGACTGGAGGGATTGAGGAAGATTAGAGAAGTTGTCATTGACTTGGCCTTGAGG
CAAAACAGGGGTTTATTTATTCAGCAAATCAATATGTATTGAGTGCCTAGAATTTAGGAATCATTGGAAGGGCATACTG
GGAAGAGATTATATTGAGATCAAAGATCAAAGAGAAAAAAATGCAAGGAAGTTTTTAGAATGTTGAGTCACAGAAGGG
TCTTAAAGCAAAGAACTATAGGAGGAAAGAGGCAGTTGAGGAATGTAAACTAGGAATATGGATCAGGGGCATTTCATGG
ATTTTACTCATTTTTATACAAGTAATTGGGATACTTCCTTCAGTCAGCGGCAACCATGAAAGGTTTTGAGTTGAGTGAA
ATGAACAGAGCATGATTTATTGGTGGATATGTTGGAGGGAAGGAAAAATTATACTGAGGAAGACCAATTAGAAAAATTT
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Fig. 6 127

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CACAATAGTTTAGGCAAACAATGAGGGCAGACAGACCCAGGATAGTGGCTGTAGAAACAAAGGAACAGATGTGAAGAAT
TTTATACAGATGTGTGTTGAGATTGCACAGTGGTGAAGGCAGGGCTTTGGGGGCACCCATTACTCAAATAATGTACATT
GTATCCAGTGTGAAAGGAAAATAAATCTTGGGGCCCCCAAAACATGAAGCTAAAGGAAAAAGTCAAGTTGAGAGACAGA
AAAGATAGAAGAAAAAGGCAAAGAGAGCTCAAAGGTTATCAGCCTGACAGTGATAAGTGATTAACAGAGGAAGAGTGTC
TGGGTCCATGAAATAGTGATGTATTTGGGCAAATAGAGTCTATGTTGTCTGGGGTGTAAGCTAAAAGAATCATTTAGCA
AATGGAGAAGTGAAATAAGAGAGAAGTATTTCAGGGTCCTCTGTTTGGAGGTGAATGACAGTTGAGGCCATAAGAGGAC
ATGATCGTATCAGGCATAGAATATAGAACAGAGATTCTCAGAATTTTCCAACTCATTCACAATACCCAGAACAGCAGAG
GACACGAAAGTTTTAACTCAAGTTATTTGATGGCACAGTTCAAAATGCAACATTTCTTTAGTATCATGTGTTTTGCTTTT
AAAAATATGGATAGATTCTTTATTCCACTGCATGATGTAGTCACACATTTCTTCCTGATTCCAGCACTAATGGACAAGG
TTTTGCTGTATTGAGATTCATTTATTTAATCAACAAATCTTTATTGAGTGTCCAGCATGCATCAGACACCATTCCAGGC
CCTGGGGATATGTCTGTGACTGAAACTGGCAAAGCCTGTGATGACGCTTACAGTGTTGTGAGAGGGAGACAGACAAGAG
ACCAGTAATGGATGAGATATTTCAGAAAACAAGTACTATGATGGAAGTAAAACATAGTGATGAGATGATGAGTGGTGGG
GGGCAACAGCAAAGACATGGAGTCAATCTAGGTGCCTATTGACACCTAGGTACATATACACCATGTGGTAAATATATAC
CATGGAATATTATGCAATCATAAAAAAGAACAAAGTCATGTCCTTTCAGCAACATGGATGTAGCTAGTGGCCATTATCC
TAAGTGAATTAAAGCAGGAACAGAAAACCAAATACCTCATGTTCTCACTTATAATGGGAGCTAAACCCTGGTTATACAC
AGATATAAAGATGGAAACAATAGACACTAGAGACTCCAAGAGGGAAGAGGGGAGGGAGTGGAGCAAGGGTTAAGAAACTA
TCTATTGGGTACAATGTTCACTATTTAGGTGACAGGTTCAATAGAAGAGCAAACCTCAACATCATGCAACATATCCATG
TAACAAACCTACACATGTGCCTCCTGAATCTAAAATTAAAAAGAAAAAAAGAGAGAAGGAAAAAAATGGCGGGGGAATG
AGAGGGTATGACTAAACTTCTCTGAAGAGGTAACCTTTGAACTGAGCCCTGAATAACAACGAGGATACAGTTATACTTT
GGGACAAGTCACCTGGACAGAGAAAAATTTGCGTGTGTAAAGGCTTTTAGGTGTGAGAAGCTCATTAGTATGGAGAAAAA
AGAAGTCTACAGAAACATGAAGCAGAATGAAATTGAAGAGTGAGCACGGTTCTGGGATACCATGCTGAGAAGTAGATTT
TATTTTAAATGCCACAGCAAATCATTGAGGGTTTTCAGCAAGAAGATTAAACGATCTGATTTAAGTTTGTAAATGATTC
CTCTGCCTAGGGTGTGGAAAATGTGTTGCATGGGGCTAGAAGTGAAAACTGGGAAAACAGCTGAGGGGATGGCTAATGG
ACTTATCATGGTGTTGAAAGTGATGAGCAAAAGACTAACCCCTAAATATTATTCATCGAGTTTATCATTTTCCTGTATA
ATTGGACTGAGTTAAAGACAGCTAATCACATACATTCGGTTCAGTATCACTACACTTTTTCTTGTCTTGGTGTTTCCTC
ATTCTCAGGACTCAAATTAATTAAGTACATGTGAGCACTTTAAAGATTTTATGATCTGTGCATGAGGAATGATGTGTGA
AGATAAGGCCAGAATCATAAATGGAAGGATGATGCCATGTTGTCAGGCTGCTCTTGCCAAAAATGTGAATGTGTCTATC
TTACTGCTGAGTCATTGGTGCACTTGTCTACTTGTCTGGGTGGACTAAAAGAGTCTCTAGTCCACTGTTTTTGTTGCTG
TTGTTGTTGTTTGTTTTAAACATTCCCATGGAAATATTTAGCAAGGACATTTTCAATGAAGAAATATGAATAGTGATC
TTTGTTTCGTGACTAATGCTGTAATTTTTGCACATGTAAATGAACTGAGCCTCAAATTATTCACATTCTGATATATCTC
TATTCAACTCTCAAACTATCATTTCATGTTATAATTTTGCTACCATTTATAAAAAACCATCACCACATTACCTTTTTAAT
AGTTGAATGACAGTGTTCTGACTTTGAGGGAAAAATCTGCTAACTAGACAATCCCATGTGACCCACAATTTGTTGTTAA
GTATAAAAAATAAATTAACTCATAATATTTAGGATTTGTTTGTTTGTTGTTTCTGAGATAGGGTCTCACTGTATTGCCC
AGGCTGGAGTGTAGTGGTGCAATCTTGTCTCACTGGGTTCAACTGATTGAATGCAGTGAGACTCCACCTCCTGGGTTTA
AGTGATTAATATTTAGGGTTTTTTTTAAATCATGATGTCTTTGTTACTTACATGGAAATAAACTAATTTATGGTTGAAT
TAGAAAATTCTAACCATGAGTACTATTTATATTTTTCAGGAAGAATACATCTAGTAGTGGCACAAGCTTTCTAAGTCTC
TTCTCATTATATTAGAATTTTACCTAAAATAATAAATATTAAAAATATGGTATATTGTAAAATTTTCTAGCTTTACTAA
ATCATTTTACATGCTGAGAAAAATCAAGAAAAAAAGTTGTTTTGAGTGGTCTCTTGGTAAATGCAAAACACTTTAAAGT
TAAGACTATGTCTGCTTTACTTGGAGTTGAATCCCCAGGATCTAGCATAGTGGTTGGTATTCAAGAAGCTCAAAACTTT
GGGATGAATAAATAAAATCATTAGTTGCTGGAGTAGTCGGGGGCATAATGTCTTTGATTCAAACTATTGCTTCCAAAGG
TTTGCTTAACATAGATGTGAGTCTAACATTTTTCTTCTAATGCTCACATCTTGCTGCAAACGTTTTCATAAGTTCTACT
TTCAAGTTCTTTGGCAATATCTTGACAGACTACTTTATAAGTGAGGCCAACTAATAAAGTAGTTACATCATTTACATCT
GTATCTTCAGTCTAATAATTGGTATAAATCCAATTTATCTTCATATTCTTGATTATTCAACAATGCTTTGTCCTGTAAA
AATACACCTCAAAAATACATCTCAAGTGCCTCTTTTTGAATCAGAACAAATCTAATCTAACTTGCCCACACAGAGCTCC
ACCAAACTATCCCTTGCCAGGAGTTCTACCTTCCTTGCTGCTTTTGAAACAAACAAACCTTGATGATACTTCAAGCAGT
TTTTTTTTTCTCTGCTTGTTTGTGCGTTTGAAGCATTGCTCATAACCAAGATTTTGCTCTCTTGCCTTTTAACTTCAAA
CTGTTTTTCCAAATTGATTTATATGTTATATATTCAATGGATGTTTCCGTGGAACAATTTTTTCAGAGTTTTTCCACTGAA
CAGCTAAATAAGGCTCATTTTGTATCTCCACCGTGATTTAAAAACCTCTGTTTTAAATGAACATCAAAATATGACTTAC
AAATTTGTGGCCAACTGCAGTGCTCATTTGGAGTTCCATAATTGTGCTTGGGGACTGTTACATGATGCCTTAAATCACC
AAATGCCCTGTGATGTACCCCATGCTTCAAGGTTGTTCTGAATTCATCTCAGTATCTTTTCCATTTTTACCACTTACAA
GCTTTTGCTCCAAACATAACTGCTTGTGACCACCTCTGTAAATTTCTAATTGAAACTGTAATTTTAAATAAATACATCT
TTGAAACAATAAATACAGAGTGACAGTTCTCTAAAGTTAGAATCTCTAGACTTATTGCTATTTTCAGACTAGCTTCAAC
AAGAAATGCAACTAAGCATAGTTTCATGTCTGTACACAGAACTTTTGTTTAAGTACATTTGTGGTACTTGAACCACATG
TGGTTGAACCAGTGGTTATGTGTGTGCATGTTCAATCAAAAATAAAATAAACAATTATGGTTTGTAAGAAGACACACTG
TTTTATAAATTCTCTCTCTCTTTCTCTCTCCCTCTCACTTATAGTTGAATTAGAAAATTCTAACCATGAATACTATTTA
TTTTTTAAGGAAAAATCCATCTAGGTAGTAGTATAAGTTTTCTTAGTCTTTTCTCATTATATTACAGTTTTAGCTAAAA
TAGTGAATATTAAAAATATTGTATTTTGTAAAATTTTTTAGTTTTACTAAGACATTGTACATGTTGAGGAAAATCAAGA
GCAAAATTCTCTGACCCATTTTTTAGAGAAATACTAGCCCTATGGTTGGGAATTATTAGTGATATCTGTGGTCACAGTC
GTGCATACGTTCACTTTGATTTGATTACAGTGTTAGTGACACTTGCTTCCCCAACAACTCTGATGTACCATATTTTCCA
GTCACTCATTATATGTGAATATTTTCTACTTACAGGACTTAAAACATGAAGTTTGCTGACCTGGAATAGTATGAATTAT
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Fig. 6

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TCTGCTTCTAAGGTTGTTTCAGCCAGCCTTATAATGCAACTAGTGTGAGAAGCAGTGACACAAAAAAGAATAGAGCCAA
ACTTCAGGTAACACCCTTTTAGGAGTGGAGAGAAGGGGACCTGTAGAAGTAGAGAAAGAAATCTGAAAAGGGGAAAACT
AGGAGCATCAGGGTGATTGATGCATGTGAAGCAGGAAGAAAGGCTTAAAAGACATTGTTAGTGGGGAAATGAAGACAGA
TAAGAAAAGAAACAGGCCTTTGGAGCTGAGGTCTAGGAAATTATTGGTGGGGGGAATGCAACTGAAATTGCTAGGAGTT
GAGTCAGGCAGCTATGAGCAAGTTCAATGGTAGGTATAAACCATGCTTTCAAACAAAAAAGTTGAAATAAGAAAACAAG
AGCCCAGATGCTAAGGACATGGGGAAATTAAGAAAACAGTTGAGGTAGAGAAAGAATTTAAGGAGGGCTGAATGCTATA
ATAGCTGCCATTATTTGACTGCTTATTATGCACCGTGGAGAGTTCTAAATGCTTTAGCTAATTTAATCTTCATGAGACA
CCTAAGAGATAAATCCTGCTATTATTATTCATATTTTGGATAGGAGATTAAGGCACAGAGAGGTCGTACAAACAAGTGA
GTGTCAGGATTCAAAGAAGGAGGTTTTGCTCCAGAGGCCTCCTGAATTTAACCACTGTGCTAGAGACACAGGGACATTC
AATGGAGCATTAATGTCCCGAGCTGGTGACAGGTCGAGCTGAGGCTATAGATGTGTTGAGGAGTCAGCATCAGTGGAGG
AGGAAAGAGATGCGGATGGAGGGAGACTCGGTGTCTTTGAAATAGATAAGAGAGAGAGACAATGTAATGGTTTCCACTT
GGCACAGAAAGAAATGCCTTACTAATTAGAGGACCAGGGAGACACAAGAGGAGCCTAAGAAGAATGGAATGAATTAGAA
TCATCACTGTGGAGTCTGTGATAGGGAGTCAACTAAAGTGGAAATAAGTAATTACCAAATAACAATGAGGGGTCAGCAT
CGGCAAAGGAATATGTTTGTAATAGAGGCTTCGCAGCTGTATCTGGAGCCTGGGAGCGCAGGCTGAGAGGGACATTGAG
ATGTAATGAGGGATGTGACATCTTAGACAGAGAGAGAGCTGTCTCAAGAACTTCATTGATATCCCTGTCTGTA
ATAGAAAGAGGTAAGTATAGCAAAAAAGTTAGAGCCATAATTGGTTTTCACATTTTAAAAAATTGGATTATGATTAAAGG
AAAACATTTAAAGACCTTGTCTAAAGGAATATAAAATTCTTGAGTCTAGGCACTCTCCATAACTTAAATTAACTTCCCC
ATTTCTTCTACAGGAATACTCCTCTGTTATAAGGATTCAATAAATATGGTGACTTTATTGTTATTCGAAAGATTATAGG
GAACTTTAGGAGGTGGTATGGGTACAGAGGAGTCCATCTACAAAAACGTTTTTAGCCCATCTTCTAAGCTAAGCTGATT
TTGGAAAAGGAGCTCTTTTAAGGAGAATAAAGCTACAAAAATGACCTTTAAAACTGTCCTATCCAGACTCTGGCTCTCT
AAATTGGAATCTTTGACAAAACCACAGCTGGAAGACAGCAGCCAGTAATACAGCCATTAATACAGCTCCATGTCCA
AAGAGGAGGGATGTTTGTTTCTACTAATTACACAAAAACATGCAATATTCAGCCAGGTGTGATGGCTCACACCTGCAAT
CCTTGCACTTTGGGAGATCAAGTCAGAAGAATCACTTGGGCCTAGGAGTTTGAGATTAGCCTACACAATGTAGCAAGGC
TCCATGTCTTTTAAAAAATAAAATTAAGGTAAAAATAAAATAAAAATGCAATATTTACTGCTAAGATGTGCTTCTTGAG
GATGGACTTGAAATGGTCCACCACCAGGACATTAAAGACAGGGTCTGCTCCTGCTTTCTCCTGAGAGAGGAAGAACACT
GGTTAGATTTACAGAAGTATCAATACCAATATTCTTAGTGTATTTAGTATTCCTGCATTTATAACAGAGCATCCGGTAA
TTGACTTGAAATTTGACCTCAGCTTTTTCATCCAGTATGCTAATGATGGTATAAAATTTACACTCCTTACATAAATAAAT
GTACCCTCAAATGTGTCATTTTCCTGCATTGACAGATGGATCCCATGCTTAAAAATTACCACAAATTCTTATTCTGAGCA
GCTTTGCACAGCACTCTGCATCCACTTTTGTTGTTTGCCACGTTATAGCACTGCTTGTGCCCTGAGCTTGCGTCTCAAG
GTGGCTACAGCGTGACATTTTCCTTGAAGAGCGATAGGCAGAATTTAATTGATTTCATTTTGCTTGGATTCTCAAAGGC
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CTAGGGACCAATGAGGTTTTGTGTAGGAGGGCTAACTTTTATTTTCCCCTTATAGTTCCTTTGTCTGGATTTTCCCCCT
TTCCTATCTATCTTTATAAAGAGACCTTAAATGAAGGCTACAGCTATAAGATGAACAAATAGCTGGCTATTAAAAATCT
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CTGCATTCAACTGGTGACAGGACTGGGCTGAGCTGGGCTGGGCTGGGTTGGTCTGGGCTGGAAATTCCAGGAAGCCTTC
ACTCCTATCTGTGGTGCCTCAGGGCTCCTCTGTATATTATCTTTTTTCTTTCACTGTCATAGTGTTCTCTAGAGAAATA
GAACCAATAGGCTGTGTGTGTGTATATGTATACATACAGCCTGTATGTTTGTGTGTGTATGTGTGTATACATAAACATA
CAAATGAGGCATGCCTCATTTTATTGTGTTTTGCTTTATGGTGCTTGACAGATATTTTATTTTTTCACCAATTGAAGGT
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CTATCATAATTGTGTTAGGGTTTCAAAAAACCATGCCTATATAAGATTGTGAACTTAAGTGATAGATTGTGTGTTTTCTG
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CAATATTAAAATTAGGCCAGTTAATAATCCTACAATGGCCTCTACATGTTCAAGAGTTTTCAATCTTTTTCTTTAAATT
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GCAGGTTGTTCAGTTTCCATGTAGTTGTGTGGTTTTGAATGAGTTTCTTAATCTTGAGTTCCAGTTTGACTGTGATGTG
GTCTGAGAGACTGTTATGATTTCAGTTCATTTGCATTTGCTGAGGAGTGTTTTACTTCCAATTATGTGATTGATTTTAG
AGTAAGTTGTCTTGTGGCACCAAGAAGAATGTATATTCTATTGTTTTTGGGTGGAGAGTTCTGCAGATACCTATCAGGT
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Fig. 6.129

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TCACTTGACCTAGAGCTGAGTTCAGGTCCTGAATATCCTTGTTAATTTTCTGTCTTGATTATCTGCCTAATATTGACAG
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GTACTAGCAGTACTTGATTTGATCAGAGACCATTTTTTACTCTGAAAACCTTCTGCCATCTGGAGGGGTTGAGAATGAG
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TAGGATTTTCCTGAGTATGTCTTCTGCTCCATGTGGTATTGATTAGGTAGGTCACTTACGTGGCTACATTTAGTCAATA
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Fig. 6.130

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GCTGGTCTGGGCTGGATGGTCCAAGAAAGTTTCAGTCATATACCTGGCACCTCTGAGCTTTTCCATATGGCCTCTATGT
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CCTGGTTTGCAAACTGTCCTTTTGTATGCTCAATAAACTTTAAAAAATTTTTCTAACTTGATCTGATTTTAACACATTC
TATCTACTTTAGTGCTTTTTATATCCTAAACACATGACGCTAACATTCTATATAGCTCACTTGTTTATACTGTTTATTT
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Fig. 6. 13j

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TCTCTTCCCCTACACTAGTGAAATTCTATGAAGGCAGGAATGTTTGTCATTGTCACACTACTGTATTTCCAGTGCCTAA
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CATTATTCCCCCCAAATATTCATTCATTCTTGGGTTTCATGAATAGAACTGGGTTTAGAATAGAAAAGGTAATGATTTG
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ACATATAAGCAAGATCATACAGCATCTGGCTGTCTGTGTCTGGTTTATTTCAGTTAGCATAATGTCCTCCAGGTTTATC
CATGTTTTCTCAAATGGCAGGATTTGTTTTTCTTTTTACTGTTGAATAATATTCCATTGTATATACACCACAGTTTCTC
TATTAATTCATTTGTCAATGGACACTTAGGTTGGTTCCATACCTTGGCTATTGTCAATAATGCTGCAATGAAGACGACA
AGTGCAGACATCTCTTCAGCATGCTCCTTTCATTTCCTTTGGATATATACCCAAACATTAGAGTGCTGGATCATATGGT
ATCTCTATTTTTAATTTTTTGAGGAATCTTCATACTGTTTTCCATAATGGAGTTACCAATTTACGTTTTCACCGCAGTG
CTACAAGCGTTCCCAATTCTCCACATAGTCACCAACACTTTTTATGACCATCCTAATGGGTGTGAGAAGATACCGCATT
```

Fig. 6.132

GTGGTTTTGATGTACATTTCCCTGGTGATTAATGATGGTGAACACCATTTTATATACCTTTTGGCCATTTGTATGTCAT
CTTTGGAGAAATGTCTATTCAAGTCCCTTGCCTATTTTTAAAATGGGTTATTGGAGTGTTTGCTATTGAGTTGTAGGAT
TTTTTTGTGTATTTTGAACATATCAGGTACACTCAGCCCTTTGTATCTGTGGGTTATGCATCAAGGGATTCAATTATCT
GAGAATCAAAAATATTTGAAAAAACATAATAAAACAATAAAAATAATTAAAAAGTAATATAGTATAATTATTTACATAG
TGTTTACATAGTATTAGTATTATAAATAACATAGAGATGATTTAAAGTATGCAGGAGGATGTATGTAGGTTATATGCAA
ATACTATGCTATTTTATATGCAGGACTTGAGCATCTGAAGATTTTGGTTTCTGCAGAGGGTGGGAAAGGTTGAACCAAT
CCCCCATGGATACCAAGGTAAAACTATGTATGGTTTGCAAATATTTTCTATCATCCATATGTAGAAAATATTTCATTTT
GCTGATTGTTTCCTTTGCTGTGCAAAAGCTTTTAAGTCTGATGTAGTCCCATTTATTTTTTATTTTGTTGCCTACACTT
TTGGTATCATATTAAAATAATAATCACCAAGACCAGTGTCAAGAAGCTTTCCACCTGTTTTCTTCCAGGAGTTTTGTGG
TTTAATGTTTTAATTTCTTTTAAGCACATTTTTGTATTGTATGTAATAAAAATATGGATACAATTTCATTCTTTTGCATA
TGAACATCCAGTTTTCCCAGGAACATTTATTGAAGAGACTATTATTTCCCCATTGTATATTATTCATGCCCTTGTCAAA
GACTAACTGATTATATATGCAGGGTTTATTTCTGGGATCTCTATTCTGTTCCATTGTTCTGTGTGTCTGTTTTTATGCC
AGTACCTCACTGTTTTGATTACTATAGCTTTGTAATATAGTTTGGAATTAGGGAGTATGACGCTTCCAACTTTGTTCTT
CTTCTTCTTCTTTTTTTTTTTTTTTTTTTTTAGATGGAGTCTCGCTCTGTTGCCCAGGCTGGAGTGCAGTGGCATGATCT
TGGCTCACTGCAACCTCTGCCTCCCAGGTTCAAGCAATTCTCTGCCTCAGCCTCTCTAGTAGCTGGGATTACAAGCACC
TGCCACCACGTTCGGCTAATTTTTGTATTTTTAGTAGAGATAGGATTTCACCATCTTGGCCAGGCTGGTCTTGAACTCC
TGACCTCATGATCCACCCACCTAGGCCTCCCAAAGTGCTGGGATTACAGGCGTGAGTCACCACGCCCAGCCATTCTTCT
TTCTTAGGACTGCTTTGGGTATTCAGGGTCTTTTTTGGTTTCATATGAATTTTACAATGGTTTTCTCTATTTCTGTGAA
AAATTCCATTGAAATTTTGATAAGGATTGCATTTAATATGTAGATCCCATATGGACATTTTAACAATATGAAATCTTTC
AATCCATAAATATGGGATATATTTTCATTTTCATTAATGTCTTTAAGTGTAGAGATCTTTCACCTCCTTGGGTAAATTC
ATTCCTAAGTATTTTGTTCTTTTAGATATTATTAAATGGGATTGTTTTTCTTAATTTATTTTACAGATAGTTTATTGTT
AGTATATAGAAGTGCAACTAGTTGCTGTATGTTGATTTTATATCTTGTAAATTTGCTGAATTATTTTCTTAATTATGAC
AGTTTTTTAGTGAAGTCTTTAGAGTTTTTTTTTTAAATATAGAAAATGTCATCTGCAAATCTAATTTGGATGCCTTTTAT
TTCTTTTTCTTGCCTAATTGCTCTGGCTAGGGCTTTCATTACTGTGTKGAACACAAATGGCAAGAGTAAGCATTCTTGT
TTTGTTTCAGATTTTAGAGAAGCAGCTTTCAGTTTTCCACCATTTAGTGTGATGTTAGCTGTGGGCTTCTCAAATATGG
CGTTTATTGTCTTGAGGTTCATTCCTTCTATTCCTAATTGTTGAGAGTTTTTGTCATGAAAGGATGTTGGATTTTGTC
AAACACTGTTTCTGCCTCTATGGAGATGATTATAGATCATATGATCTTTATCTTTCATTGTGTTCATTCGGTATATTAC
ATTTTTTGATTTGTGTATATTGGATCATACTTGCAAATCTAGGATAAATTCCACTTAATCATGGTGAATTATATTTTTA
ATGTATTGTCAAATTCAGTTTGCTAGTATTTTGTTTAGGACTTTTGCATTTATGTTCACCAGGGATACTGCCCTGTAAT
TTTTTTTCTTATAGGGTCCTTATCTGATTCTGGTCTGTTGGTAATGCTGGCCTCATAAAATGAATTTGGAAGTGTTCCC
TTCTCTTCACTTTTTTTGGAAGAGTTTGAGAAGCATTTGTATTAATTCTTAAAATGCTTGCTACAATTCACCCTTGTTG
CCACATGGTCCTGTGCTTTTCTTTGTTAGGAGTTTTTTTTTTTAATTATGGATTCAATCTTCTTACTTGCTATTGGTCT
GTTGAGGTTCATGTAATATATTTCCCTATCCCTTTGCTTTCAGCCTATGTGTATCCTTAAGGCTTAAGTGAGTCTCTTA
TTGTCCCAATATGAGACAGCATTTTGTTGGATTTTTTTTAATTCATTCAGCCACTTTGTGTCTTTTGATTGAATAATTT
AATCCATTTATATCCAAGATTATTATTTATAGGTAAGGAGTTATTACTGCCATTTAAAAAAATTGCTTTCTGATGGGTTT
GTGGTTCTTGTGCTTTTTTTTGTTGTTGCTGTCTTGTGATTTGTTGATATTTTGTAGTGGTGTGCTTTGATTCCTTGTT
GTCTTTTCTGGATCTATTAGAGTTTTTTTTTTTTTTCCTTTGTAATTACTATGGGACTTTCAAAAAACATCTTGTAGC
ATCTTTTAATTATAATATTTTATTTTAAGCTGATGACAAGTTAATTTCAATCACATACAAAAACTCTACACTTTTACTT
TCCCCCGTCATTTTGTGCTACTGATGTCATGCTTTGCATCTTTTTGTATTGCATATCCATTAACAAATTACTGTAACTA
TGGTTTATTTTTAATTATTTTACCTTTTAACTTTTATTCTAAATTTAGAAATAATTTATACATCACCATTACCACATGC
AAGTATACAGAATTTAATTATGTATTTACCTTTTCCAGTGAGTTGTATACTTATATATGTATTTATGTTGTTATTTAGC
AGCTTTTCATTCCAACTTGAAGAATCACATTAGTATTTCTTATAAGGCAGGTCTTGTGGTAATGAACATCCTCAGTTTT
TGTTTGTCTGGGAATGTCTTTACCTCTCCATTTCTGAAGGCTAACTTTGCAGGGTACAGTATTCTTATTTGACAGGCTT
TTTTCTTTCAATGTTTTGAATATATTATTCCATATTATTTATTTGTTATCCATGTAGGCTTGCATTGGTGTCTGTGAAT
TTGAAGAAGCAAACAGTCTTTCGGTCTTTACAGGCTGATTTTAACAACCTTCTCTTCCACCAATGGCAGACCTGTTATT
AGGTATGCAGATAGGCGTGGTTCCCTCTGGGTTTCTGGAGGACTCCCCCTGGCTCTCTGAGTATGTCTATGGGTAGGGA
GAACCGTCCCCAGATCAACATGAGAGAGCTTGGAACTAAGTCACTGCTGCTTCAGGGTCCACATCTGAGAGGACTTGCC
TCCAGGGAGTTGGATGGGCATACATCTCTGGGGACAAGATTGACCTTGGGCCAAGTCTAAGTGGGAATGGAGCAAAGTT
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CAGGTCGCTAAGTGGAGACAGGACTGCTCTCAGGCCACAGTTTCAGTTTTAGGATATCATAGTTAGCAGACAGCTCTAT
ACAAAACTCTACTCCTTTATATCTCTGCCCATTTATGTTATCAATGTCACATATTACAGCTTTTTATATTTTGCATTTG
TTAACATAGTTTTATATTTATATTTTATAATTAAGTTCTATACCAGAGTTAAAAATAATTTATGCACCACCATTGCAAT
ATTATACGATTTTTATTTGTCTAAATATTTACCTTTTCCTTTGTGTTGCTGTCTAGCATACTTTGTCTCAACTCAAAGG
ACTCCTTTTAGCAATTCTTGTAAGTCAAGTCTAGTGCTAAAGAACTACCTTAGCCATTGTTTATCTTTATTTCTCCTCC
ATTTTCAAAATACAGTTTTGCCAGAAAGAGTATGCTTGGTTGACAGACTTTTTCTTTCAGCACTTTGAATATATTGTCC
CACTCCCTTCTGGCCTATAATGTTTCTCCTGAGAAATCTGCTAATGTCATTGAAGGTCGCTTGCACATGATGAGCCATT
TTCTCTTCCTGCTTTCAGGATTCCCTTTTTGTCTTTGATATTTGACAGTTTGATTATAATTTATGTTGGTATGTTGGTT
TTTCCACGTTTGAGTTTTATCAGTGTCTTGAATTTGGTTGTTCATTTCCTTCCTCAGATTTGGCAAGTATCTAGCCATT
GTTTTTTTTCAAATAAGCTTTCTGTCCTTCTCTCTTTCTATCATCTCCTTCTCAGATTTGCATTATGTGTATGTTGGAAC
ACTTGATGGTGTCCCATATGTTTCTTAGTCTTTCTTCACTTTTCTTCATTCTTCTTTTTTTTCCTCTTTGTATCTCTGGC

Fig. 6.133

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TTCATAATTTTAAGTGGCCTCTCTTTGAGTTCATTTATTCTTTCTTCTGCTTGATGGAGTCATCTGTTGAAATGCTCTA
GTGAACTTTTCAACTCAATTACTGTATTCATCTCCATGAATGTTTGTTTTCTTTTTGAAGT: .TTGATATTCTCATTT
TGTTCCTGCATCATTTTCCTAATTTCATTTAGTTTTCTGTCTGTGCTCTCTTCTAGCACATTGAACTTCTTTAATATTA
TTTTTAATTATTTGCCAAGTAACATAAATTTTTATTTCTTTCTGGTTGGTACCTGGAGGTGTATTTTGTTCCTTTGATT
GTGTCATGTTTCCCTGTTTCTTTGTGTGCCTAGTTATTTTTTTCTGTGATTTGTGCATTTGAAAAAGCAGCCACCTCTT
CAGTCCTTATAAATTGGCTTCATACAGGGGAAGACTTTTACCAATTAGCCCACATAGAGATTTTGGGAGCCTCTCCAGT
TTTGTTTTGTTTTTTTTTTTTTTTACCACGGGATACATCTTCTTTGAATTTGTGTGTGTTCTTTCCCAGTTAGAGTGAT
TTGCTTCTTTTTCTTCCAAGAGCTTTTAATCTCCTTCTCCGTCTGGTGTCTGTCTGTACCACTGCAGGTTCTGTGGCAT
TGCAATAAGCCACTGAGTTACCTTTTGCTCTCTGCAGACCCCAGGCATCCAAAGTATGTAGATTCCATTAGTGCTCTGA
ATCAGGCAAGACAGAAACCAGTCTCTCAGACAACACTCTGAAAAACCAGAACATTGGACATACATTCCACTCTTCTCTT
TCTCTCCTAAGGGAGAAGTCATAAGGTGTGATTTATTTTTCCTGATGACACCAAGCTGTGCTGGCTTGGAGGAAGGGCT
GTCATGGTTGACGTGAAATGCCTTTTCTCATCTGTTTCAATGAGACTATTATTTTCTTTAAGTTTGCCTCAGGCACTGC
AACTTCTTGACTGGTTTCTAGACTTTTTCATAAAACTTTTTGGATCATATATTATTGCTAAGTTGGCGTCTCTGGTGGAG
AATTAGCTCTCTCGCTGATGAAACTCTACAGAATACTTTTACAG.ATTTAATATTGCTCTTATTTCATCTCACACGTAC
ATAGTCACTTTCCTAACACTTATATAATTGTTCTCTTCCCAATAGCTAGTAGTGTTTCCTGTAACACATATTGACACAT
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CACACTTTTGGTTGTTATTGTGGCTTCATGGTAAAATTTTATATGTATTAGAGCAAGACCGTAACCATGTTATATTTTA
TAACATGGCCTAATTCCTCCAAAAAGTAAAATAAAAAACTAAAAAAGAGATGATTTTCTTGCAATAACATTAAACTTAG
AAAATTATTCAAAGGAACATCAGTCTTGATTATATAAGTCTTGGAAAAGTTGTATAGATTTCTTAATCTAAGTTCTACT
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TCACTTTTCTTATAGTTTTATGCCTCCTATTTCTATTCCTCACATTGGGTTACTGAATTTGTTAGAGGGTTTTTGAAAAA
TGGGGGCAATGTTACATTTGGGCTAAAACTATGAACATTGATATCAGCTAAACCTGGTCTCAAAACTCTGAAGGATCCT
CAGTTAGAACTGAGAGGTCTTAAAAGACGAGGTGGCTAGAGGGAGGTCTGCACACACATTGCCATGAGTCCTCAGACCA
CATGAGACATTCTTTTTTGTCTGAAGCAAGGCTGGAGCCAGGCCTAGATGGCCTGTGGAAAGCTCCCATGGAACCAGCC
TGCTGTTCTCATTCCAGGCAAACTAACCTCTGTACAGTTGATAGTGTCACTATGGACAAACTAACCTGTTTAAGCTTCA
TTTCCAAATCTGTATCATGGGAATAGTAACAGTTTTTTGTACAGTTTTTATAAGAATCAAGTAGCTACCTATACAATAT
ATGTAATACAATAAATATTACTAAGTTTCTATTATTATCCTGATCTTAAGCATGAACCTATGCTTTACCAGTAGCTATA
ATGTTAAATATAAGTTTAATATAAACATTCTTTTGATCCATAGTTTGTTTATCAAGGGGATATTTTGTAGTTATCTGAA
TATCAAAGCCAAAAGCAAAGCAACATTTCTGGCTCTCTGCATGTGGAAAGGGAACTTAGCTTTACTTTGTCTCCCCTTT
CCTCCCACAGCCCAATATTTGTTAGTTTATTCTAGATTTACATACCCAGTTTATCATTATAAATATATCATCACTTAAA
TGATGGTTTTAGGTTTTTCTTTCATGTTTATACACATAGTTTTGATTGGTATTGGTATTTAAATAAAACTTGAGAATGA
AGACAATAGATCCTTTGTATTTCTTTTAAGGTTGTGATTTGACCCCTGGTTCCTCAAGTGAGAAACTTTTTCAAAACACT
TGAGAGAGTAAACTGGCTGAGTTTTACCTATACTTACTGTAAGACATAATTTTTAGCTGCTAAAAAAACAATGTTAGTT
TCTTTTCTACACTTGGGATTCTATTACTGTATGTAACTGTGAAGGAAATAATTGACAGAACTCATGTTTTTCCTTATTT
TTTTTCCTTCCTCTACTAACAAAATAGAATTGACTTTATTTTTTAAAGTCAGCCAGGAAATGTACAATGCTATTTACAA
AAGGTTAATTGAATCCTAGAGTTGCAGATCGAGTTTCTTGTTGATGAGAATATACAGTAGAATATTACTGAATGAGAGA
TTTTTCTATTCCTTTTTTGGAAACTTTTTGGAAGTTTTGATGACTGAGTTATAGGAAATTGGCTTTAGAAGGACTTCAG
TGATTTCTGTCTCTTCTAATCAGTATGAAATCACAGCCTATATTTGGATTATGCTCTGTGAATGTTTGAAACCTATCTG
TTATCTTTGTGTTAACCTGAGGTAGAAATTTTCATTTTTATATAGAATTAGTTTTGCAAGAGTATCTTTTCTATATACA
AATATCCATTTGAAAATTGATTGTGTTAAGTCGATACAATGCCCTTTTCTATATTAAAGGACAGGATGTTTATTGCCTT
GACTTGTTTGTTTCCTGTAAAGTTATTTGGGCAATAATAGCATTCAGTTTTAAGATAAAACATACATAATAAGTAATGA
GACAAGAGGAAAAACATTTGGCATAAGGCTTAGAGAAATTTAAGGCTCATTTTCCATTATTGCTTTAGTGCATTATTAA
CAATTTCATGGGGACATGAAAGACTAATATATGGAGAAAATGTCTGCTATTGGCATATAATATGTTAAGATTGCAAATA
TGACTTTTAATCCACTGTGATAATAGCCTATTAAATTGTATCCTAGTCCTTGAAGAGTCACTAACATTATCTTGTTTAT
ACTACAAAATTAGGTACAGGGAAACTTTACTTTTACAGAACTTAATGTTGCATCAAACTAACTGGACAGGAAAAAAAG
GAATGTGGCATTGGAAGTGATAAAACAAAACACATCCTCTGCCTATGACTGTCTTCTTCCTGGACCTATCTGACTATTT
AACATGCTGACCTTTACAATGCTCTGCATAGTGTAGGCTCACTGAAAGCACTGAGTGTCTGAAATCCTTCATTCTTATT
TAAAACAAAATAGGTGAAAATACTTAATCATGATGTCTATATATTTGAAAGAATATTTGAACATGGGGAGAGACCCAAC
CAGTTTGAGTGGCTGAAGTTTTGGAATTTTTATAGAAATTCCCTATTATGTGTTTCAACTTTATGCCTGACTTATAATA
ATAGTTACAAATACTACATTGCTAGTACACCAATATCTAATACCATTTCCTCTTTCCTAATTTCTTGCAGTTGAAGAA
ATGGAGATGAGTATGTCTATATGCTTCACTCTCTTTCCTCAGCAAGATTTATTTAGCATATGTTTTGTTCCAGGCACTGGC
TAAGTGCTAGGGGTACAAAAGTTGTTTTGTCTCCCTTCCTTCCAGATGTGTCCATGTGTCCAGGCTAACAGAGTGAGTA
GTCACACAGGCAAATCATTCTCCTACAGTTGTCAGGCTGAAGGCTGACTTGGTGTCATAAATAAAAAGGCAGAAGATAT
TTGTATTCATTGTCTTGGATGATATTTTCAACTTGAGTAGGCAGATTGAAAAGAAAAATGCGGCCGGGTGCAGTGGCTC
ACACCTGTAATCCCAGCACTTTGGGAGGCCGAGGCGGTGGATCACCTGAGGTCAAGAGTTTGAGACCAGCCTGGCCAAC
ATGATGAAACTCCGTCTCTACTAAAAATACAAAAAATTAGCTGGGTGTGGTGGCAGGTGCCTGTAATCCCAGCTACTTG
GGAGGCTGAGGCAGAAGAATCACTTGAACTTGGAAGGCGAGGGTTGCAGTGAGCTGAGATCATGCCATTGCAATCCAGC
CTGGGCAACAAGAGCAAAACTCCATCTCAAAAAAAAAAAAAAGAAAGAAAGAAAGAAAGAAAGAGAAATGGTTGAAG
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Fig. 6.134

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GATAGCAGCTTGGACAATGCCTAGGTTTTAAAGACAGTTAATTGAGGAGCCAGCAATGAGACTAAATGAAAATGACTGG
AGAAGCAGATAGACAGAAAGCAAAAATAAATGTCCAATTTATGGGGTGATCTACATCTTTTTTTGTCTGCAACAGTCCA
GGTGTTCTCTTTTGTCGATGTCACAACTGGCTAGCACTTTCTTTCAACTTGCAAAAGTGTTATGTACTCCAGTAAACC
CTAAGTTCTCTGTTACAGCTTTGATCATATACTGGTGATTATCTGTCTATTATATTTTTCCTCAAATGAACTGTAAGTG
CTTAAGCAAAAGGAGGTGCCTAATTCATCTCTGCATTTTCAGTACCCGACACAAGCATCAAAGGTAGTAGGTGCTCAAT
AAATGATTAATTGGGAAATTAATAGGAATGGAAACACAATTAAGATAGGTTGAACTCCTTTAGAGATATACACATAAAA
TAGTACTACCTGTAACAAGAAAATGCTAAATTTTTGTGGATTACCCCAACAGAAGATTATCTGTGGCTCGTGTACCCCA
AATTAAGTGTTTCTGATTGGCAAGTGTCTCTCATGTAAATAATGATTAAGAAACCATGGCTTCTTCCATCTTGTGACTC
TACCATCTTCAACACATGGTGAGGGGCTACAGAGGGGAAGAGAAATAGGGGGTTGTACTCACAAAATTTTTATGTGGCA
GGCCTGAAAGTGGCACATACACAGACCCATACACATTTCCTTAACTTATAATCAATGAGCATGTAATGTAACACTCACT
GACATCTCTATAGCTACCTGTAAGGGAGTCTGGAGATGTGGTCTCACTGCATGCCAAGAAAGAAACATTCTTTCCTGCA
ATGCCTATATATGTATATGTATAAATATATATATGTATATACATGTTTACATATGTATGTATATATATATGTGTGTATA
CATATACTGTATGTGTGTATATGTATATATACACACACACACACACATCTTAAGGGGATTTTCTTATAGAAATTTTATTTC
TCCCCCCTCTCTATATATATATGAAATAAATATCAGAGACTGTGAGATTATTAAGAACCACAGAAATTTATTTAACCCT
AGTAATGTGCATGCCTGTGTTTTAAAGGCCATTAAGTAGTTTATTGTCTACTTTTGAAAGAAAAAAAATTAAACAAAGA
AATTAATTAAATATCTTGGATAAAAACTGAAACAAACAAAAAGGAAAATAATAATAATAAATTTAACATTCATCCATTG
TAAGTGTCAATTATTATGGAATTGCTTACTTTAGGCATGTATTTCAGAAAGAATTTTATGCAGTAAAAAGCATAACATT
AAAGTTATTTGTCTGTCAATTATTCCATATAAGTTGGCTAATTTTTGAATATAATTTTATTTTACTACAAGGTAGCAGG
TTCATATTACAGTTATTTCAATATGTGAGCATTTCTTTTATTTTGATATTGCATATCTTGAAGCCGAATATATTTCTAA
GTCCCATCAATAGCAAGGTGAATGTTGTATCATTTTATTATTATGACTTTTATTATACTTTTAAAATTGAAAAACAAGA
CATCTTTATTATAGGGAATACAGAAGCTCATAATTAATAAAGATACTAAAACAAAATTATTACCAATATTGCACAGGAG
AGCCATTTTAGAGAATTAGGATACATCCTTAAACAATTGTTTTATGAATATAATATGTGACTTTTTTGTTTTGTTTTGA
TATTATTGTTTTATAAAAATGGGAGCACTCTATAATCTGTTCTTCCCACTCAACATTGTGTCATTAACATGATTACGTA
TTCTTCTACATTATTTTAACAGCCTCATAGTATTTCATCATATGATGTATCAACATTTACTAAGCCAATCCTTCAGTAC
AGCAATTAGATTACATCCAGTTATTTGCTATTATAAATGAGCTGCAGTGAACATCTTTGTGCCTGGATATTTGCCAAAC
AGTTTTATTAAACTATGTCTAAAATGTCAAAACCATTAGTTCTCAAACTTTAATATGCACTGGAATGATACTGGTGACT
TATTCAGACTCATAAGCCCTACTCATAGAGATTGTGTTTAGGAAGATCTGGAATAAGACCCGGGAGTCTGAATTTTAGC
AAGCACCATCTATGATTCTAATGCAGAGGGTTTGAATGTCACATTTTGAGAAATAATGACTTGGAGACATTAGAAATAC
TATCTTTTTTCCATCTTCTCCCTGCCATAATGCCATTTTCTAACAATAACATAAGATACTTATTGCTCTGGTGATTAGT
TTTACATCTCTCTCCAAAATAAAGGCATTTAACCTCTTACCTTCTACCTGATAAGGTTGTTCTTTTTATTTCTGAACCA
TAACTGATACACATTGATCTAGAGTGTGAAAAAAGCCTCTTATACTGTTTTGGAATGGAAAATGTTAGAATATAGCCCT
CTAGTGCTTTATCATTTTTATTGTAAAGATAAAAGTATTTATAGAAAGTGGGTTTAAACTAACAGAGTATAAGCATGAG
TGTAAACTTCATTTTTAGTAGAGATAATTATCTCAAAAAAGTTACGTCTTGAGGCAGTTTTATCTAAAAAGAATGTCA
AAAACATTAAGTCATTCAACAGATAACCTCAGTTATATAAAGCATTTTTGCAATTGCAATGCTAACACCCAGAGTAGGC
AGCTAAATGCTTATAAAAAGATGTTAACTTTGGCCAGCTATATTTAGTGCTACTGAAGCTTTGCTAGACTCATCCTTTG
TTCTTCTTTTCTATGTTCATCCAGAAGAAGAGTATTTACACATGTATTAGTCAGGGTTCTCTAGAGGGACAGAACTAAT
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GGTCTGTAAGCTGAGGAGCAAGGAAGCCAATTCGAGTCCCAAAACTTCAAAAGTAAGTAAAGTGGCAGTGCAAGGATCC
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TGTTTTCAGTCTGTGGCTCAAGGCCCAAGAGTCCCTGGAAAACCACTTGTGTAGGTCCAAGAGTCCACCTGTTTAAGAA
CAGGAGTCTGATGATCGAGAGCAGGAAACATCCAGCACAGAAGAAAGATGCAGGCCAGAAGACTCCGGCATTCCAGTCC
TTCCATGTTTCTCTGCCTGCTTTTAACCCAGCTGTGCTTGCAGCTGATTAGATGGTGCCTACCCACATGGAGGGTGGGT
CTGACTCTCCCTGTACACTGACTGAAATGTTAATCTTCTTTGGTAACACCCTCACAGACACACCCAGGAACAATACTTT
GTATCCTTCAATCCAATCAAGTTGACACTCAGTATTAACCATCACAACACATAAGTCATTAACTTAATCATGACCTGTT
TCCAATCCAGGAATGAAGTTCTGTTCTTAGAGGAACTTCCTAAGTACATAAGTACACAGAGGCTCACTTTCCCAATCTA
AAAGCTAAGACTGTTCCAAGTGCATATGCTTTCTGTTTTTTTCTATATTTGTTATCTCACAGAAACATTGATTTGCCAT
TGGTTCTTGCAAAATCAGTATTTTATAACTGGAGGTGGTGTTAAGGTTTATTTTGTATAACCTTTTGATTTTATCCCTA
TGGAGACTAAAGACCAGCAAGGTTAAGACCAGCAGATCTGGAACAGGTTATTATAGAAATTTGTATATTATGTTGTTGC
TAGCAAATGGAATTATTGCAAGGAGATGATGGAAAAGATAGTATTTCTGATATCTCCAAGCCATTGTTTTTCAAAATGT
GTTGTTCAAACCCTCCGCATTGGGATCAGAGGTGGTGCATACTAAAATTCAGACTCCCAGGTCCTATTCCAGATCTGCT
GGTCTTAGCCTTGCTTGTCTTTAGTCTCTGCAGACAAAAAATCAAGACATTATACAAAATAAACCTTAAGACCCCTTCC
AGTTATGAAGTACTCTGATTTTGTAACAACCAATGGCACCTCTTAAGACCACCTGAGTAATTACTGGCAGAGCAGGGGA
TAAAATTTCTATCCTCTCATTTAATGACAGATTGTCATTCATTTCTAATGTTTATGTAGGGCTGCTCTCATCTCAGTTC
TGTTTAATTTTTCAAATGAAAATGTGAAGCTTATTGAGTTATGATCAATTAGTGTGTGTTCTGTTATTTTCTACTAATA
CTCACTCTGCCAGTGACTATACTTGTTAAGGTAGTGCAGCATGGTGGCTTTTAGCTCCAGTTTTAACTAGAAGAGATTT
GGGTTTGGATCCTGACATGTAGGGTGATCAATCATCCTGGTTTCCTTGGGACTGTGGGGTTTCCCTGGACATAGGACTT
TGAGTGCTAAAATCAAAAGGTCCCAGGCGAAGTAAGATGATTAATCACCCCACTAGCCATGTGGCTCTGTCATGATAGG
TCTCTGGGTTTTTTAATGGGTAAAATTGGAAAAGCAAGAAGAAGTAACTCATAGGATTTTAGTAAAAATTAAATGAGAC
AGTATATGTAAAACACGGCACTACTTAATAAAGACTCAGTAAGTATTAGTCATTATTATTGATGTTTATAAAGCAAGT
AGATTCCTCAAAGCACAGATTCTCAGAATTTAGGATTTGAAATAAATGAACTTTCTTTTTAGCTTTGACCTCTTTTTTT
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Fig. 6.135

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TTGTTTGTTTTTTAAATCTTCTCAGTACCCTTCTGTCGTCTTGTCTATACAACTGGACAATACAACCCAAGGTTGGATA
GAGTCCACCATCTCACATCCTTTTACCTTTTTTTCAAGATATAAATCAGGAAGACAAAAATTTTCTATTTGGGAATCAAT
GTGAAGAGTGAGCATCAAAGACAGCATAGAGCTATGGAGAAAGGGGAAATTAGCCTGGATTATAATAACAGCTGCCATT
TTGTTGAGTGCAAACTCTTTGCCAGATATTGGTGCTTCAAATACTTTATATATTTGGTCCACACAATGACTCCGAGAGG
TAAATATTATTATCTCAGTCTTCAGATGGAGAAACAACCTCAATGAGCCTAAGTGACTTGCCCAAAGATAGTTGTATAG
TAAGGAGTGTAATAAATGAGGTCTGGATGTTCCAAGCTCGATGAATGTCCAACTTTCAGCCAGCTGGTTAATTTTGAGA
AATATAAACTTCTTCACCTGTAAAACTCAGAAGGGAAGGAAATTGCCATATATGAGTGCTAATCCCAATGCAAGTAATG
AACTATGCACATCATATGCATGATCTCATTTAATTCTTGTATGTAAATATTAGTCCCATTTTTATAATAAAGAAACAGA
AGCCTTAGAGAGACTAAGCAACACACTATCAATACAAATAGTAAAGTGGCAGTGCAAAGATTCAAAAACCCAACACTCT
ATAGAAGCATGTTATAACATTTGGTGATTCTATAATTCTCAGTATGTGGATATAATAAATATTGCTAGAGTAGACTTTC
CAGCCATTTGGAAATAATTATTGCTCATGTGATAGTATATTTTTCTCAACTAGAATCAAAATATTAACTTTTGACCTGG
GTTTACCTTTGCATGTGGATAACATGGATCAAATTTGCAAAATTCTGCTGAGCTGTGACTTAAATACAGTTCTACTGGG
TCCTCAAGTTTCTTTTAAAGTCTTTTTTTTTCTTACTACCCTCTCAGTCTTATAAAAATTGGCAAGAGTATATCTGAGTA
AATGTGACATATGAAAACATATGTTTCAAATATGGCCTTAAATATGCAGAAAAAATAGGACTTTTCTTCCTCTCTCTTC
ATGGGCATGCATGTAAGAAAGATATATATAAATATACAGTCATGTGCTATATAACATTTTAGTCAACTATGGACCATAC
ATACCATGGTGGCCATAAGATTATAATACTGTATTTTTACTGTATCTTTCCTATGTTTAGATATACTTAGATACACAAA
TATTTCCCATTGTGTTATAATTGCCTACAGTATTTATTACAGTAACAATGTACCATCTAGGTTTGTTTAAATACACTGT
GATGTTCACACAAGGAAATTTCCTAATGATGCATTTCTCAGAATGGATCATTATCAAGTGATGCATAACTGTATGTGTG
TGTGTGTATATATATATATATATATATATATGACACATGCATATGTCAATGTATATATTTAATGATGTTTGCAAT
TCATTTCATAGAAGTTATAAAAACATTTTGATTTATCCTTGATTTTATAGAATTCACACACAAAGCTAATAAATTCTAA
GATTACTGAAGATGTTAGTTAACTGTTTTAACCAAGCCACACCTGGTGAACAAATTTGCTATTTTAAAAATAAATTTAG
ATTTTAGTTAATCAAATATTGTTCAGTTAATGTCACTGTTTTCTAGATAGCATTTTTCATGTATACATTTTACTTAATT
TCTTTCCCAGGAAAGTATTAAGCCATTATGTAATGATAAACAGTGTGATTTTTGTTGATTATATGTACTTTCTTGAATT
ATTACCTCAGGCCTCTGAGGAAACACTACATTCCAGTAATGAAGAGGAAGACCCTTTCCGCGGAATGGAACCCTATCTT
GTCCGGAGACTTTCATGTCGCAATATTCAGCTTCCCCCCTCTCGCCTTCAGCACAGTTGGAACAAGCTGACTTGAAAAGTG
AATCAGAGAACATTCAACGACCAACCAGCCTCCCCCCTGAAGATTCTGCCGCTGATTGCTATCACTTCTGCAGAATCCAG
TGGGTGAGTGCCCTCAGATGTCATTTCCCCATTTTATATTTTAGATGGTGATTGTTGTCTGTGGTCTTTTTGAGTTTTTG
TGGCTCATTGCTTGATTTGAGTGGGAGAACAATGGGGTTTCTAGAGTCAAAAAAAAAAGGATGAATTTCTCAGGTGACA
TTTCTATATCTGCAGGAGAAGACATTTTAGTGAAATTGTTTTACATGGAAGAGCTGAAAATGTATTGGTGAATGCTAAT
AAACATTTGCCAAAGGGCAGTTGTACTTGGAAAATTAATATTGGCCATATGTGACTTGATAGAAGACATTTTAACAAAT
GGAATCAAGAATAAATTGTATTGTAATTAGAATCAAAAGAAATATGCAATGGTATTCATGAGTCTAAAAATTTTGCCTT
TGACATTTTGAGGTATTGCTTACCAGAATACTATTTCAGCTCCCAGTATAAGGTGATTTTATTTAGCATTAGTACTTAT
AGAGGAGTCAGCCACAGCCACGTCTCCATCATGTAACATTTTTATTTCCTACTTTGCAGGGTATAATTTTTGTTTTTAT
TCTTTTTCGTGTTGTTTACCAATGGAGTTTTACACTTTAAAGCAATGTCACCAGTTTAAAAATTGTGAAAGAAAGTAAG
GATATTGGGCACATCAATTATTTGTTCTTGTGACCAGCTGCATTTTTAAGAAGCCTGGATAGAAAGGAAAGAGGATGAA
GCCAGTGGGCCATACATAGATAGACTCTTGAACATTTGCTAAAACTACAAGAACAAAACAATTACTATGCTACAAATGT
GATGGTCCCACATTCTCTGCCAAGTTAAAGGCATCTCTGGGGAAAATGTCTTTTGGATCTTGTTAAAGTTAGGAAGGTT
GTGCTCAGAGGAAATTTGGCTAGTAACTCAACTCAAAGATACTGAATTTAGCTGTTTTACTTGTTATTGACCAGTCTAA
GGGACCTGTGTTAGAAAAAGGTATGAGGAAAATACTACTAGATAAATAATAAGAAAACAAATATTCACCTTGAAGCAAT
GAAAAGAGGAAATATATTTTGTTTCTGTTGTTGTAATTGAACTATGTGTTTAAAGAAGAATGCACACACATTGATCAGA
CTCATACTAAAGCAGATGGGAATTTTATATGACAGCTTGACCTGAACATTTTTTGAAAAATGCTGTTCCCCTGAAACTA
TTTCTCCTTTCTTTGGAAATAACATTGCTGCCGTTCATTCTGTGTAGAAGAGATTCACAGATATCAGGTGTATCCAGGG
ATCCCAAGATCACCCTCGGGTTCAATGATTTGCTAGCAGAGCTCTCATAATTCAGCAAATAGTCATATTCATGGCTCTG
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TAAGCTTCCAAGAGTCCCCAAGTGAAGTAAGACAAGATACACTTAATTCCTCCAGTAATGAGTTGAGACAACACTTGTG
AAATGTGTTCACCAGGGATGCCTTTACCAGGGAAGCCTATTAAAGACTCAGTACCTAGAGTTTTTACTGGAGGCCAGTC
ACTTAGGTACCCTCTGCCTAGGATATACAAACTTCCAGATTTCCAGAAGGAAAGTGGATATGCAGCATAAACCACATTA
TTTGCACAGACAGTTTAGGCAAATTGAGCCAGTCTTACCATTTAGAGAATGGTGGATCACTCCTGAAAGTCAGGTCTGC
TAAAGGCCAATGTTTCAAACAGGTCTTCCTATATTTAATTGACACAGTCCTGATTAACTGTTTACTACACACCAGGTTA
CCCTGATATATTCAATTAGCTTTGGTATCTCAAAAATTTTGGTTGCTAACCATGATTTCTCATTTTGAGTTGGTGTTAT
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CTTGAGCTGGGCGCAGTGGCTCATGCCTGTAATCCCAGCACTTTGGGAGGCCGAGGCAGGGATTGATTGACCTCATGAA
TTTGAGGCAAGCCTGGGCAACATGGCAAGACCTTGTCTCTATCAAAAACACAAATAAATAACTGGGTGTAGTGGCATGC
GCCTGTGGTCCCAGCTACTCAGCAGGCTGATGCAGGAGGATCGCTTCAGCCTGCTTGAAGGCAGAGGTTGCAGTGAGTC
AAGATCCTGCCACTGCACCCTGGCCTGGGTGTCAGAGAGAGAATGTATTGCTAAAAACAGTTTCCAAGTGCCAAGGAAC
TTCAGGGTTCCGAATTGTCTATGTCCCAGCATTTCTGTTGATTCACATTCATTTCCATCTTCATTAAGATTTAAGAATC
AAGTACAATTGACCCTTGAACAATACAGGTTTGAACTGCACAGGTTCCTTTATATGAGGATTCTTTTTCAACCAAACACA
AATGGAAAATATAGTATTCCTGGGATGTGAAACTCATGTATACAGATGGGCTGACTTTTCATATATGTGGGTTCCACAG
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Fig. 6.136

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GGCCAATTGCAAGAATTGAGTATGCATGAATTTTGGTACATGTGGTAGACCCGAAACCAATCCCCTACATATACCAAGG
AATGACTATGTAAGCAATGTCATGCTTACTATTTACAAATTTATCCAATATTATGAAGAAAATAATTTTTCTTTTTGAA
AGATGGAACAAGGGTTTCATTTCAAAAGAAACAATAGACAATAATGGCCAATCAACAAGAACCAAACTAAGTAGATATT
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CAGCACTTTGGGAGGCTGAGGCGGGCAGATCACAAGGTGAGGAGATCAAGACCATCCTGGCTAACACGGTGAAACCCAG
TCTGTACGAAAAATACAAAAAAAAATTAGCCGGGCTTGGCAGTGTGTGCCTGCAGTCCCAGCTACTCAGGAGGCTGAGG
CAAGAGGATGGCGTGAACCCGGGAGGCAGAGGTTGCAGTGAGCCGAGATTACGCCACTGCACTCCAGCCTGGGCGACAG
AGTGAGACTCTGTCTCAAAATAAAAAACAAAAAACAAAAAACAAAAAACGAAAGGTGAAGTTCGGGCTTATATTCTGTA
GCCATTTTAGTTAACCAGGAGCCCCAGATTCCTTTACAGCTGTCAGAAAGAGCTAATGGGAAATTTAGCCTATTTTAAC
TGAGTTGTCTCCAGCCCATTGCCTCATTATTAAAGGTTAGTATGCCTTTAAGTCCCATTGCCAACCTATACTGAGACCA
TACCTTAGCAATTGCAAATCTTTGATGGTCAAGGCGACTTTTCATGCCCTCCAGATTTATTTTCCTTTCTTTCTTAACA
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CATAATGTATTTATTTTTACTCTCTTACTAATGAACCTTTGGGTTGTTTCTAGTTTGAGGGGATTATGAATGAAGCTGC
TATGTACACTTATACCAGTCTTTTTGTGAATATATGCACTCATTTCTTTTGGGTATATTCCCAGGAGTAGCATTTCTGA
GTCATAAGGAGGCAAATATTTATTTTCATTACTGCCCAACAGTTTTCTAAAGTGTTGTACTCTTTTACCCTCCCACCAG
CAATACGTAGTAGGTACAGTTGTTCTACATTCTCACCAACACTTAGTATTTTCAGTCTTTTTCATTTTGGTCATTCTGG
TGAATGTGTAGTGGTATACTACTGTGGTTTTAATTTACATTTCCCGATGAGTAGGAATATTGCTTACCTTTTGATGTG
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CCCCATTTTTCACATATTGTGAATACTTAATATATAACATCAAAAACTGCAACTCCTATGCTGGGAAAATATTAGAACA
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TGACATTTCTGTTCAGAAGTCTGTATCTGGCATCTTCCCGCACACATGTCTTGCTTTCAAGTGCTGTTCAGTTATGTTT
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AATCCTTAGGAGGAATTAGGATCCTCCTTTCAGCCTCTATCTTGGGCACTGACTTACATTCCCACATGTGGTCCATGTGCC
CCTAGACAATCTAAGGAGAGTCTAGCTTGCCACACATGTGGAAATCATTCCCAGTTCACAGGTGGGAAGTTATGGGAAT
GAAGAAAATGTGGGGGAGGTAGCAATAAAACTCTCTACTTACATTTCATTGCTGCATCTGCTATAATAAAATGAGAGCC
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GTTACAAACACTTTCTCATCCCTAGTATCCTATTACCTGTCTGAAGAAGTACACTGTGTACCTTCGTATCACTATGTCC
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GTACCTTCATGTTTGATGTATGTTCAAGCTTTAAGTGATTTATCTAAGTTCTTATTGAATGTTATATGGAAGATCTATA
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CTTTGTAAATTTGTGATTATTTTTGTTTTGTTTTGTTTTGTTTTTGAAACGGAGTCTTGCTGTGTCACCCAGGC
TGGAGTGAAATGGCGTGATCTCGGCTCACTGCAACCTCCGCCCCCCAGGTTCAAGCGATTCTCCTGCCTCAGCCTCCCA
AGTAGCTGAGACGACAGGCATGTCCCACCACACCAGGCTAATTTTTGTATTTTTAGTAGAAATGGGGTTTTGCCATGTT
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Fig. 6. 137.

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GGCCAGGCTGTTCTCAAACTCCTGGCCTTATGAAATCTTCCTGCCTCAGCCTCCCAAAGTGCTGGGACTACAGGCGTGA
GCCATCATGCCTGGCCCCAAATTTGTGATTTTCA__ATTAAAATTAGATTCTAAGAGTGATGGAGCACTTTAAAATATC
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CTGCGTTCACCAACATCAGAGCATCTAGTCCCTATTTTCAAAACTTTCTTGTGCCAATATGACTTGTTTTAGTAATAGA
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ATAAAGTTTATAATACATAAATACTATACTTCTTTACCATTCACTTTCTAGAACTTTCCCAGCTTTTGATCACATATTT
CAAGGGTGGTCTCCCTCTTTACTCCTCTACTAACTCTTCTGTGTCTCAGTTAACATGTGATTTTGGCTTTCTTATTTCT
CCAACATTATTTAATTCAGACTAACTTTATTTCTTGAAGGTATATACCAGACATACCTCGGAGATATTGTGGGTTCAGA
TCCAGGCCACTGCAATAAAGCAAATATCACAATAAAGCAAGTCACACATACTTTTTGGCTTTCCAGTGCATATAAAACT
TATGTTTACACTATATTGTAGTGTGTAATAGCATTGTCTCTAAGAAAACAATGTACATGCCTTAAGTAAGAAATACTTA
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CAATGTTCATGACTGCTGACTGATCAAGGTGGTGGTTGCTGAAGATTGGGGTGGCTGTGACAATTCTTAGAATAACAC
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CCTTTTTTTTTTCCAGCTTTTATTTTAGGTTCAGAGGGTATTTATGCAGCTTTGTTAAATGGGTAGCTTGTGCACCCTGT
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GCATCTGCTTGGCTTTTGTGAGGCTTCAGGAAGCTTTTATTCATGGCAGAAGGTGAAGGGGGAGCAGCTTTGTCAGATG
ACAAGAGAGGGAATAAGAGAGGGGGGCAGGTGCCATATTCTTTAAACAATTAGATCTCACAATAACTCATGACCACAGGG
AGGGCACCAAGCCATTTATGAGGGCTCTGACCCCATGACCCAAGCATCTCCCACTAGGCCCACCATCAACATTGGGAAT
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Fig. 6.138

CAAATCTCAACATGAGATTTGGAGGGTATAAATACCCAAACCATATCAAGCAAAGCACAATAANACAAGTATGCCTGTA
TTTATACACATATTACAAAACAGAATTAAAAACAATAAAATCATACATGCAAGGGATTCTGGTAAATATACTGATCCCC
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GGGCACGTGCAAGAAAATAGACCACTATAATCTTGTTTGTTTCATATAATGTACAAATATGACTGCAGACAAATGGAGA
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TTCCAGGAAGAGAGAACAGCAGGTCAAAGTCCAGGGATGAGATTAAAGTAGAGTGTCTGCACTTAGCTAAATAAAAACT
GCTAAAACATGGCTAGACCAGAATGCCAATTGAAGGTACAGTTGATAAAACTTACCATTAACTTTTCATGTAACCAATA
TGTGAGGTTTTGTAATATTTTCCAGTATGCTAAGCAGCTATGATATAAACACTGAAAAGGCAAACAGTTGGATTATTCC
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ATCCACCACAGGGAAGTGACATCTTTAGCGAAGAGTCAAATTTGAGAGAGGTTAGGATCTGGGATGACTTTTTGGTGGC
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CCTCTGACTGATGGGTGTTAAACCAGGGAACAAGAAGTTGAGCATAGCTAGTAGGTCTTTTGGAAAGACCATAACGTTT
GTCCTAGTCCCAGTAAGTAGGTCTGCAATGGTGTCCTTGGGTTCCTAGTACATTTCAGCATCTATTGGAGAATAGGAGC
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TAGAAACTAAAGTACAGAAACTGCCCTGACTTTGCTAGAAGTTTAGGTGTAAAATTAGTAGTAAAGTTAGTAAGGAAAA
ACCAAAGGGGAAGATTGAATTTTATTTTAATGCATTATTTTTTTTACATAAAAAGATAACAAAAGGAGGGTAGTGAAGAG
AAGAGTCTCTTGAGATGTCCGTGGCAAAGTTTTTAAAGCTCTCAATAAGGAGACAAGGACTGGGCTTCATTGCAATTAT

Fig. 6.137

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TCCCTTTTGACTGTTGAAGCTGGTAGGCTTACAATGGACTCCTAATGAGAGTGGTTTGCACTACAATTAATTAAGTGCA
TGACATTATAAACTGCATGTCTCTTCCCTCTGGAGAATATTCTTCCTATATAAGTGCTATTTTGTTTTAATTTCTCTCT
CTGTCTCTTCCTCTCTGTTTCCCTCTCTTTGTCTCTCTCTCTTTTTTATTTATTTATTTATTTTTTATTATTATACTTT
AAGTTTTAGGGTACATGTGTACATTGTGCAGGTTAGTTACGTATGTATACATGTGCCATGCTGGTGCGCTGCACCCACT
AACTCGTCATCTAGCATTAGGTATATCTCCCAAAGCTATCCCTCCCCCCTCCCCCCACCCCACAACAGTCCCCAGAGTG
TGATGTTCCCCTTCCTGTGTCCATGTGATCTCATTGTTCAATTCCCACCTATAAGTGAGAATATGCGGTGTTTGGTTTT
TTGTTCTTGCGATAGTTTACTGAGAATGATGATTTCCAATTTCATCCATGTCCCTACAAAGGACATGACCTCATCATTT
TTTATGGCCACATAGTATTCCATGGTGTATATGTGCCACATTTTCTTAATCCAGTCTATCATTGTTGGACATTTGGGTT
GGTTCCAAGTCTTTGCTATTGTGAATAGTGCCACAATAAACATACGTGTGCATGTGTCTTTATAGCAGCATGATTTATA
GTCCTTTGGGTATATACCCAGTAATGGGATGGCTGGGTCAAATGGTATTTCTAGTTCTAGATCCCTGAGGAATCGCCAC
ACTGACTTCCACAATGGTTGAACTAGTTTACAGTCCCACCAACAGTGTAAAAGTGTTCCTATTTCTCCACATCCTCTCC
AGCACCTGTTGCTTCCTGACTTTTTAATGATCACCATTCTAACTGGTGTGAGATGGTATCTCATTGTGGTTTTGATTTG
CATTTCTCTGATGGCCAGTGATGGTGAGCATTTTTTCATGTGTTTTTTGGCTGCACAAATGTCTTCTTTTGAGAAGTGT
CTGTTCATGTCCTTCACCACTTTTTGATGGGGTTGTTTGTTTTTTTCTTGTAAATTTGTTTAAGTTCATTGTAGATTCT
GGATATTAGCCCTTTGTCAGATGAGTAGGTTGCGAAAATTTTCTCCCATTTTGTAGGTTGCCTGTTCACTCTGATGATA
GTTTCTTTTGCTGTACAGAAGCTCTTTAGTTTAATTAGATCCCATTTGTCAATTTTGGCTTTTGTTGCCATTGCTTTTG
GTGTTTTAGACATGAAGTCCTTGCCCATGCCTGTGTCCTGAATGGTATTGCCTAGGTTTTCTTCTAGGGTTTTTATGGT
TTTAGGTCTAACGTTTAAGTCTTTAATCAAAAGTTGTCCCTGAAGGAAAATTAACTCTGCTGTTAGTCCACTGAGAAAG
CTTCTCTTTTAATACTGGAACATATCTTTGTCTAACTCTCCCTCACACCTAGTGCTAAATTTATCAGCGTTGTTCCTGA
TACAAACATTTTGAAGACAATCGTGAGGAAAAAGATGAAAAAAGTTGCATTTAACTTTTGTTACAGGTGTCCTTGTTCT
CCAGAAGTTACTGTCTTTGTTGTAGTGTGCATGCAGTCACAGGAGTGTGCATACATGTGTGAAGTACTTGAGAAGTGTA
TTTGTCCGTTCTCACACTGCTATAAAGAACTGCCCTAGACGGGGTAATTTACAAAAGAAAGAGGTTTAATTGACTCACA
GTTCAGCATGGCTGGGGAGGCCTCAGGAAACTTACAATCATGGCAGAAGGGGAAGCAAACCCTTCTTCACGTGGTGGCA
GCAAGGAGAACTGCAGAGCAAAGTGGGGAAAAGCCTCTATAAAACCATCAGCTCTCATGAGAACTCACTATCACGAGAA
CAGTATGGAGGTAACTGTCCCCATGGTTCAATTACCTTCTACCAGGTCTCTCCTGTGACACATGGGGATTATGAGAACT
ACAATTCAAAAGATGAGATATGGGTGGGGACACAGCCAAACCATGTCAAGAAGTATTTCAGAGACAAGTCATTCCCCAA
ACCTTATCATCAGGGTCAAGACCTGGCAGGTGAGACATTAGCAGGTGAACACAACATTCTCAGAAACCCTTGAGAGTAC
TCCTTTCCCTCCAGGCATCTTCTTTCAACCTTAATATATTCTATTCAATTATATTCACAGCTTTTTACCTCAATTATAA
TTCCATTCTTCCAAATTTGTTTTCTTCTCCCATCTTTCCTCCTAATAAAATATCTATTCTTCAGATAGTCTTCCTTTGT
AAAATAGAAAGGATCTACAAAAGCCATTCTTTGTGAGATGTGAATCATTTCAAATAAATTAATGCCTTGTTCTAATGTG
AATGGTGGACTTGTGAAGAGAAGTGGGTAGATATTTGAATTTCTGAATGCTTTAGAATATTAGTTGCACATGCAGTAAT
ATTTCCTGTAGCTTAGAAGAAATTGGTTGGCTTAACAGAAAATGCAATTAAAATAGTTTACAAATAGGTTCTGGCATCA
ATAATAGAAATATTAAAGAAACTACATATTATATCAAGATGCTTCTCTTTTTTTTCGGGGGGGGGGGTGCCTTTTCTTCCT
TGATGAACACAGTACTCTTTCTTCTTTGTCTTTTATCTTCCATCTCAAATTGTCAATAATAAATTTGAAGAATGGAGAA
TAATACATTATCACGTGCCAGACCTTATGCTAAACACTGGATATGTTATATTGTGTTTCATACTCAGGATGATGCTACG
GTAGGTACCATTATTATCTTATTTTACTTATAAGAAAACAGAAGCCTGGAGAAGTTAAACAATTTCCTCAGTATCAGAA
AGAACCAAGATCAAATATCTAGGTTAAGGTATTTTATCTTAACCTAGATATGCCAGAATATCAAATCTAGGTTTCACTA
TTATTTTGTATTCTATCCAAATTTCTAAAACTGCTAATGATGGAGGGCTGTTATATGGTTCTAGCTTTATATATTTTTT
TATTTCAACTTTTATTTTAGATATATGGGTATTTGTACAGATTTGTTACATGGGATTATTGCATGATGCTTAGGTATGG
TATATCCCATTACCCTGATAGTGAGCATAGTACAAGATAGGTAATTTTTTAATGCATCCCACCCTCCCTCTACCCTCTA
GTAGTCCATGGTGTCTATTGTTCCCATATTTATACATATGTCCACGTGTGCTGAATGCTTAGCTCTCACTTATAAGTGA
GAATGTGCAATATTTGGTTTTCCATTCCTGTGTTAATTTGCTTAAGAATATGGCACCCAGTGGGCCGGGCGCAGTGGCT
CACACCTGTAATCCCAGCACTTTGGGAGGTCGAGACAGGTGGATCACCTGAGGTCAGGAGTTCTAAACCAGCCTGACCA
ACATGGAGAAACCCCGTCTCTACTAAAACTACAAAATTAGCCAGGCGTGGTGGCGCATGCCTGTAATCCCAGCTACTCG
GGAGGCTAAAGTAGGAGAATTGCTTGAACCCAGGAGGTGGAGGTTGTGGTGAGCTGAGATCGGGTCATTGCACTCCAGC
CTGGGCAATAAGAGTGAAACTCACTCTCAAAAAAAAAAAAAAAAAAAAAAAAAAAAARGCATATGGCTCCCAGCTCCATTCAT
GTTGCTGCAAAAGACATGATTTTATTCTGTTTTGGGGTTGCATAGTATTCTATAGTATATATGTACCACATTTTCTTTA
TGCAATCTACTATTGATGGGCACCTGGGTTGATTCCACATCTTTGCTATTGTGAATAGTGCAGTGATGAGCATATGAGT
GCATGTGTCTTTTTAGTAGAATTATTTATTTTTGTGGAAGTATATACCTGGTAATGGGATTGCTGGGTCAAATGGTAAT
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TAAGCATTCCCTTTTCAGCCTCGCCAGTATCTGTCATTTTTTGACTTTTTTATAATAGCCATCCTGACTGGTGTGAGAT
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GTATGTCTTCTGTTCATGAAGCCCTTTGCCCACTTTTTAATGGGGTTATGTGTTTTTTGCTTATTGATTTGTTTAAGTT
CCCTGTAGATAGTGAAAATTAGGCCATTGTCATATGCATAGTTTGCAAATATATTCTTCCATTTTGCAGGTCTTCTGTT
TACTCTGTTAGTTTCTTTGCTGTGCAGAAGCTATCTAGTTGGCCATCTTGGCATCTAATCTTTAATTTTCTTTCTAAAA
TATTTGAATGTTTTGCAATTTCTGACTGAAGTTACTTCCCTCTTTCTGAAGAAGGCCCTGCTGACATCAATAATTATCT
GAGAGTGACATAAGCTGACTCCGATTATGCCAAAGTAACCCTTACGTGGTATGAAAAGAAAATGAAGGTGACTATGATT
CCTGGGGCATTTGAGATTCTGAGAAAACTCCAGGTCAGCTCGCCATAAAAATTCCCCACACCTGTAGTTTAATTTACCA
ATTAGGAAATTATATATTTGGGGCTAAATGGTTTCTGATTTGTTGGCCTGAAAGAAATAATGTACTTTTAACAACTATA
AAAGTTCTCTGCTAGGAATATCTTAAATACAGTGAAAATCTGCCTTGACAGTGGACAGTAAGTTAGATTTCATTGTTGT
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Fig. 6. 140

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AAATTGGCACCATTAACTAAAACCCTTGTCCATTCAAAATAATATGTATTGTCACATAACACTAAGCTCTGTTATTAAA
CTGCCACAGATTATTGCTTTAACAGTTTGAAAATGATTTTTAGAAAATTGACTTTGTACAATGCTATATTAGAGTCTGT
TAATTGATGAGAAAACCAAACTCAGATGACATTTGCAAAAATAAAATTGGCATCAGTTGAAACATCTGTACCATTTCTC
TTATTTAAAAGACATTTTTCATCTGGAAAATGACTACTTCGATTAGCCTCAGATCACAACCACAAAATAAAGTTGTCTA
TGGGTGGCTTCTTATTTGCATTTAAATTATTATGTTGCAGCATTTTGTATGCACAAGCAAATGGAAGATTGTTTTGAAT
GACATAGGCCTATACACCTGAGAGAGAAATATGTCATCTACAAAATGTAAAAATATAGGTTTTTGTAAATGCCTCTGAA
AGGATTATAAGAATCTACATTTTCCCCTCCCTCCTGGAGTTTGTCTTTCTATTAAGGCTCTAGACGAAGAAAAGCTCTA
ATATACTTTCCTTGGAGGGGTCATGAGACAATGAAGCATAATAGGATTTGAAGTATATTGGATATAAAATTCATGTGAT
ATCCTCAAGTAGAGAGATGTTATTTTACTGTTCTTTAATCTTTACTGTTGGCGATAATTGCCTTTGTAAACTAACTTTA
TGTTGTAAACTAGTAGCGAACCAGAGATTGAGTTGAGAGGTGGGGGTAACAGATGAGATCCTCACTTGTGAGGGCTCCA
AGGGCCTGTGAGGCAATGGAAAGGGCTGACCTTGCTACCTGGATGGGAACAATGCTGTTATCATTTTCATGTGCTCCTG
TTTAGAAGATGAGTTAACATCCATAAAAACAAGTTAGACTAAGGGTTGTCACACTACAGCCTGTGGGCCAAATGCTGCC
TGCCACTTGTTTTATAAATGAAGGTTTCTTGGAGCTTGCACTACTACTGCAGAGTAGAGTAGTGGAGATGGAGACCATT
CGGCCTGCAGAGTCAACAATGTTTAATATCTGGTGCTTTACAGAAAAACTTTGCCAAGCCTTGATTTAGACTTAGTAAC
TTTAGTAAGAAGTTACTGACTGCACAAATTACCAGGATATTTAAGCTTGTAAAGTATTAAAATACCATAACTATCTGCT
TCTTGTGGCAGGGAGGGGGCCAAGGAGAGGTGAAGTGGTTATAATGCTCGTGCATTGGTTCCTTCATTCACCAGAGTGA
TTGTGCGCCCTCTCTGAAACAGGGATAATGCCAGGTACTCAAGATACAGCAGTGAGCAAAAAGAGGCATAGTGCCTTCC
TTAATGGAGATATGCATTGGAGTCACATAGCATAATAATGTTGACAAATATCCCTGCTCTCATGGATCAAGAGGTTAGA
GTTTTAGGGTAACCATGGAAGATCTCACTGAGAATGTAACTTTGAATAAAGGATTAGAGGACTTGAGGAAGCCAACCAT
GCTGCTAGAGTTTCTCTGCAGAGGAAAGGGGAGTGATGAGATATAGGTGAAGGACAATGTGAGCCAAGATTGTTTTTTT
TAAGTTGAGGAATTAACAGCATGGTTGTATGCTGAGGGGGACAATTCAGTGGAGAGGGAAAAATTGATGTTGCAGGAAA
TTGCTGGGATAATGTCTTTGAGCAGATAAAAAGGGATTGATCCAGTGTACAACAGAAGGGACTAGTCTTGGTTTTGGAT
CAAGGACAGTTTATCCATAGTAAAATAAGAGAACGTAGAGTATATGGGCACAGATGCAGGTCATTGGGTAGTCGTGGTG
GTTGAAAATTGTAGAAGTTCTTTTTTTTAAAAACTTTAATTTTTCTTAGTATGTGCAAAGGCCTTGAAGCAAAATAGAGCA
TGGTATAATCAAAGAATTGAGAGGAAGAAAAACATAACTCAACTGTAAATTAAGCAATACCATGGAGCCAGATGAGGCTG
ACGAGGTACACAGGATACAGACGTCAGCAAGACTAAGAATATGGTAATGCTAAACCAATTAAGAATGGGGTTAAATCAA
ATTTGCCTTTTTAAAAAAAGATGACTGTGGCTACTGATAGAGATTGGATAGGAGAGTGCAACAAGAGTTTCTGTGGAGA
GACCAATCAGGAAGCTACTGCAGTAATTCAGGTGGGAAATGAAGGCWTCATGGTCTAGAGTAGAGGGAGTGGGGAGAAG
AGAAATGATCAGATTTTGAGAGATATTTAGAGGCAGAGTAGATAGACTTACCTCACTTATGTTTGGGTATAGAAGACAT
TGCCTATGTAAAGACTGAGTAGAGAAGAGAGCCTAGGAGTAGATGTGGACACTCAGGAGAAGGATTGAGATCCTGGTTT
AGAACACTGAGTGAAGAAGCTGGGTATTGTTAAAGAAGAGAGCCAAACTTCACAACTGCCTAGCAGTGAAGAAGCAGAT
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TTTTTTAAATCCTATAAATATATTGGTTAATTTTATTTAAAAAAATTGAACTTTTCATTCTCATAATGAGACCAAAGTA
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TGCATCTGTTCCAAATATGGGTATAACACAGACTCTGTGAGTAGCCTGCAATGTGTTTCCCTTCCACATATTGCTAGAT
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AAGAAATGACTGTAGAAGTAAAAGAACAAGAGCTGGTCTTAGGAGTTCCATCCCTTCATCAGTCTAGGAGGGTTGGGAT
CTTAGAGGCATTTTTGAGCTGAAAACTCAAGGCAGGAAACAAGCCATCCCACTCTGAATAGGGCTGTGTTTCTCACCTG
AACAGTGAGAAGAGGGTGTTATATTTTCTAATAAAATCCCACATTAAGTTAGACAAAAACTCTGTATAACTATGGAATA
GACTTTACTAATGGAAAGAGTAATGTGAAAAGCTAAATAGGAAAATTTTAGGCACGTATTTTCACAATAGATCTTGCTG
TACTTGGGGGAGGCTTGTCTTCTTGATTACTAAGGTTAACTTAGTGCTTCAGAAGCCAGCCCGTTTTTATGTTCTGCTG
TTTTCCTGATTTTGGTGTGTCTTGATGCCACTGTGTTCTAGTAAAAATTCCCTTCAATAGACTGCCAGGAGAGCAGACC
AGGGAGTGTGAGTGTTTTGTGCATGGGCGAGCACATACAGATGTTAGGTGACAGTGGTTATTCAGTCATGCTTTCAGAC
CTGGAGAACAGTTCAGGTCAAAATCTCCAAGTCTGTGGAATGAAGCTTTGGGGGAATTAATCAAAATGTTACTGAAAGA
AAAGGGAAATCAAATCTGACTTCCACCCCTCTGAGATTTTTGTTGAGGACACAGAATGAGAGCATAACCTTCAGGTCCA
GAGTGGAATATAGAGAGATCTTCCACTGACAATATGCTATCAACAAAAGCAAGGTAGAGAGGAAACGTTGAGCCTCAGC
AACCTGTAAGTCAAGACCCTCTCCATCATCCCTAGATGCTTTTGCTCTGTCATTGGGATTGTGTCCCTTTCTATAACAC
TGTGACTTCTTTGGATCTCCACAGCCTGACTGTGTTCACGTTCTCCTCTCCAGTCTTTTAGTTTCTCTTACAGAGCTGG
TGCATTTCTGTGGGGACAGAGCAGCTGGAACCTGGGCCCGAAGCTGGTTCAGATATGGAATTCCCTTTAAGAGCAAAAA
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TACATTTTCCCCTATGCTTGGCTAAATATTCTTTGATTGTCTCTTTAAATGACACTGAAATTGTAATATTTTCTATGTA
GAGATTGGGAAGATACTTTAAACTTTTCTTCAGGATTGTTGATTTTTAAAAAATTATTAATAGTTAGAAAAGTTTTGTA
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CAGGGCATTCGATGTGGTGTGTGTGTTTGTGGGTGTGTATGTTGACTAATCTTACATACTTGTATTCTTTTACCAGTTT
GTCATCGAAGTGTCACTGCATTTGTCATGAGATTTAGATCCTCTCTGTCAATGTCAACATTTTATATCAAATCAGAAAA
AAATTAAAAATTGTTTCAATGTGTGTTAGTTTGATTAACTCCTTTTCATTAATTTACTTGTCAAGCAACCCAAGGGCCA
ATTCATCATATCTATTTGAAATTTATCTCTTTCTCTTATTGAATTACCACTTTTGAGATGATCTGGAAAAAAATTGTTA
CAATGTACTGTCTGATATCTGAATAATTTCTATTGATTTAACCACTTGGCTTTACTGCATTTTCTTCATACTCCAATAA
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Fig. 6.141

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TTGTATTTGAAATTTTCTTATTCATTTATATTTTTAGTCCTTTAATCTAAAAAAATTAAGGATAAAAAAGCATCCTTCA
CGTATATTGAAATCAGGAGTCAGTAATTTTTCACTTGTTTAAACATTCAAAACATCTTTCTAACTCACTTCTTGATATT
GGGATAATTTTAATTTTACTTTTGCAAATTTTGCCAAAGCATCTGGCTAAATAAACACACTTGCCAGGGCACCGCCCCA
GAGCCTCGGCAGGTACTTATGCAAGCAGAGAGCCCTGCAGCTTTGGCTTTTATCATCTTCACCATAAACCCATCTCTGA
TTCTCCCCTTTCCCTACAGCACTTTCCACTAACAATACACATACACACCCCATATCACACCTATACTATTTATTTTTCTT
TATGCCCCTTTTCTGTAGTATTTATATCCATAGCAATTATCACCATCTAACACACTACATATTTTGCTTATTTATTTAT
TTATTGTTTATCTCTCCACACTAGAATTTAAGCTTTGTAAAGGCAGATGTGTTCCTTTCCTGAGGCTGCCATATAAATG
ACCACAAATGGGTGGCTCGAAACAGCAGAAATTTATTCTCTCTGAGTTGCAGAGAGAATTCTGCATCTCAGAATACAGT
TCAAAATCAAGGTGCTGTAGGGCTGGTTCCCTCTGGAGGCTCTGGGGGATGATTGCTTGCATGCCTTTCATAGCCGCCA
GTGGCTGCTTCCAACCCTTGGCACTCCTTGGCATGGGGCTTCATGAGGTTCAATCTCTGCCTGTGTCCTCTCGTGGCCT
TTTTTTTTTTTTTAATCTGGGTCTCTGTGTCTTCTCTTTTTCTGATCTTAAACATAATTTCTTGAGGAAATCATAAATC
ATTGGACTTGTAGCTCACCCTAATACATGGTGATCTTATCTCAAAATCTTTACCTTAATTACATCTGCAAAGATCCTTT
TTACAAGTAAGTTTACATTCTTATTCCGGGTGAACTTATCTTTTGTGGACCACAGTTCAACTCACTACAGAAGTTACAA
TTTTTCTGTTATGCTCACTGATGTGTCTCCAGCACATAGACCAGAGCCTGGAATGTAGTAAGTGCTAAATAAATATTTG
TTGAAGGAATGCATTGAGAGCCTATAGTGCCAGATGCTGAGGTAGGTACTTAGAGCACTGCAATTTAAATATTCCCTAA
CTATAATTTATCTTCTCTTTTCAATTCACATATAAATATTTTTTTAATTTTTTAAATTTTTTATTTTAGATTCAGGTGG
TACATGTGCAGGTTTGTTACATGGATATATTGCATGATGCTGAGGTTCGAGTTTCTATTGATCTTGTCACTAAGATAAT
GAACATAGTACCCAATTGGAAGTTTTTCAGCCCTTGCCTCCTCTCTCCCTCCCCTCTTTTTGAGTTCCTTGTGTCACTG
TTCCCATCGTTATGTATGTGTGTATGCAAATTTTAGCTTCCACTTATAAGTGAAAACATGCAACATTTGGTTTGCTGTT
TCTGCATTCATTCATTCATTCATCAATGGCTTATGGCTGCATCCATGTTGCTTCAAAGGATGTGATTTTGTTTTTTCT
TGTGGCTGTGTAGTATTCCACAGTGTATATGTACCATATTTTCTTTATCCAATCCACCGTTGATGGGCACCTAGGTTGC
TTCCATGTCTTTGCTATTGTAAATAGTGCTGTGATAAACATACATACAAGTGCAGGCATCTCTTTGGTAGGATGGTTTA
TTTTCCTTTGGTATTATACCCAGTAATGGGATTGCTGGGTCAAAATGGTAGGTCTAATTTTAGTTCTTTTGGAAATCTC
CAAACTGTTTTCCACAGGGTCTGAACTAATTTTCATTTCTGCAAACAGTGTACAACCATTCCCTTTTCTCTACCGTCTC
GCCAACATCCGTTATTTTTTTACTTTTCAATATAGTAGCCATTCTGACTGGTGTGAGATGGTATCTCCTTGTGGTTTTG
ATTTGCATTTCTCTGATGATTAGTGATATTGAGTATATTTTCATGTTTATTGGTTGCTTGTATGTCTTCTTTTGAAAAG
TGTCTGTTCATGACCTTTTCCCACATTTTAATGGGGTTATTTGGTTTTTTTCTTATTGATTTGTATAAGTTTCTTATGG
ATTCTGGATATTAGGCTTTTGTTGAATGCATAGTTTGCAAATATTTTCTCCCATTCTGTAGGTTGTCTGTTTACTTGGT
TGATAGTTTCTTTTGCTGTTCAGAAGCTCTTCAGTTTAATTAGATCCTAATTTTCAATTTTTGTTTTTGTATGCAATTG
CTTTTGGGGACTTAGTCATAAAATCTTTGCCTAGGCCAGTGTCCAGAAGAGTATTTTCTAGGTTTTATTCTAGGATTTT
TATAGTTTGAGGTCTCCCATTTAAGTCTTTAATCCATCCTGAGTTAAGAGGTAGGGGTCCAGTTTCATTCTTCTGCATG
TGGTTAGCCAGTTTTCTCAGCACTGTTTATTGAATAAGGAGTCCCTTCCCCCCTGCTTATTTTGTTGACATTGTCAAA
GATCAGATGGCTGTAGGCGTGCAGCTTTATTTTGGGGTTCTCTATTCTGTTTCATTGGTCTATGTGTCTATTTTTGTAC
CAGTATCATTCTGTTTTGGTTACTGTAGTTTTGCAGTATAATTTGAAGTCAGGTGATGCCTCTGGCTTTGTTCTTTTTC
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AATAACATTGATAACTTCATAGAAATACTGTTGAATCTGTACATTGCTTTGGGCAGTATGGACATTTTAGTGATATTAG
TCATTCAAAGCTATGAGCATGAAATGTTTTTCCATTTGTTTATGTTGTCTATCATGTGTCATTTATTTCAGTAGTGTTT
TGTAGTTCTCCTTGTAGAGATCTTTCACCTTCTTGATAGTCCTAGGTATTTGTGTGTGTGTGTGTGCACCTGCGTGT
GTGTGTGTCTATTGTTAAGTGAGATTGTGTTCTTGATGTAGGTCTCAGTTTGAATGTCACTGGTATATAGAAATGCTAC
TGATTTTCGTACATTCATTTTGTATCCTGAAACTTTACTGAAGTCATTATCAGGTCTAGGAGCCTTTTGGTGGAGACAT
TAGGTTTGTCTAGGTAAAGGATCACATCGTCAGCAAAGAGAAATAATTTTACTTCCTCTTTTCCTATTTGGGTGCCTTC
ATTTCTTTCTTTTGCCTGATCGCTCTGACTAGGACTTCCGTTACTATGTTGAATAGGAATGGTAGGAATGGGCATCCTT
GTCTTGTTCCCATTCTTAAGGGGAATGTTTCCAGCGTTTGCCTGTTCAGCATGATGTTGGCTTGGGGTTTTTCATAGAT
GACTCTTATTATTTTGGGATATATTCCTTTGATGCCTAGTTTGCTGAGGGTTTTTATCATGAAGGGATGTTGGATTTTG
TCCAATGCTTTTTCTGTATCTATTAAGATGATAAATATGGTTTTTGTTTATAATTCTGCTTATCAATTCACTATAAGTTC
TTTGAAGATGAAAACTCTGTCTTTTATGGTGTTTGTACATTTTGAATTTCTTAATACTGTGTATGTAGCTGACAACACA
TAAATAGATATGCAAATGAGTAAGAGAGTCATTTGGGATCACTGCTGATTTAACTTATCTTGCCATGAAAGCATTCTGT
TCCTCTTTCCCATGCTTGGTAATATATCCTTTTGGAAGAAACACTCTTGCCAAAAGAGAATGGCATTATTACATCCAAG
GGAAATATTATTTGCCCAAGAGTGTAAAAATGTATGAAAGAAATATGTAAATAAAATCCTATATGTAAAATCTAATAAT
ATCAAAGTTTAATGATGAAATTTGTCAAGATGATTATAAAATATCACAGTTAAACTGTATATTATATTCATGTTTCTTG
CCATTTGTATGCTAAAAACTTATAATGATTTGAATGTACAAAATCAAAAATCGGGAGAAAAGAAGATAGATCAGAGCCT
ATTACCAATAATTTATCCTTTACCTCCATAAATGTTTGCTAGAAATGCCAAAGGCTCTAATTTGTCTATTTAAATGTAC
TTCAAAAAGCATTCCCCAAGTTGCCTATTTACTCTGTTATCTCTATAAATCAGGAAACTCTCCAGGCTTTCCACTGTG
AATGTTGTATGGAATTGATAGAACAATTTCTTTTTTATGTTGTACTCATTCACCATTTAGTTACTTAGACAATAAACTT
AGAATGACAAATTCTAAGTCTCAAAATATGAATGAAAATTCACTCATTTTCAGAACAAAATTTAGCATCACACTCTCTT
GAGCAAAATCTCACTAGAAAAAAAGATAGAGAAAATACGTTATAGCATTTGAAGTCTGCTACCTCATGTCCTTTTATCT
GCTTTCTAATTTATTAGAAATAAATATGCAATTAAAATATGACTTTCCTTAACCTTTACATAATTATAATTATTATGAA
AAATATGTGTAAAAATATACATGTTATATTCCTTTTCTCTATACTACCAACAGCCACTCCCATTCATGTACTCATATTA
GAAGGGAAGAGCTTTGAATATCATATATAACATTTTCAGATAAAAGAACTTAGGTCCAGAAAAGCTAACTCTTTTGCTG
GTGACTAAACATACACATACGTGCACTGACACACACATATTAAGTTACTGAGTTGATAGCAAAAAAAAAAAGTGCAACT
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Fig. 6.142

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TTATATGGAAGTAGAATTATGTAAAAATGTCTGTCTAATTCTATACTGGAAAATATCTCATATGAAATGGCTCCTTCAT
ATGACAACACTCAGTATTCAAAAGCAAAGCACAACAACTAAGAAAGTTACTTTCCCCAACCAAAAAAAAATAATAATAAT
TAAAAAAAGAAATGTACCTATTGTGGACTAGCTGATTGGATACCATGGCCACAGCCCTGATTTTCAGTAAAGACATAGT
TTCAAGAATGTCATTCAGATGGTTTCTTAAGTAGGGATGTTGTTATTTTTAGCAAAAGAAGCCTTTTGTCCTTGTCACA
TCAAAACACAAAGAATTCCATAAGTCAGTGCCAGTCTTGTTTTTAATGCCTAAATATCTGGCCACATATATATTTATGT
TATGTAAATGTTATTGCCTTGTTTTTCAGAATAAGAATTTAGAACATTAGAAATTGACTTTTATAATGCAATTTTTTTC
AAACTCTTTTTGGCTTTTCTCTAAATCTCAAAATTTGTTTCAATTTAGGCAATGGAATTGGACTAGTCAGTCTAATTTA
TTTTTGTTAGCCAGTTTCTCTCAGATTCCAGGGAACAAAGCCAAACTATAAGATATCTACTGAAAACATTTGTAAG
AAATAATTAAAAGCAGACTCTGAATAAGCATATAAAATGAAGCATTTTTATCACAAGCTTTATAAATTTTTTAAAAACT
GTGTTCATTTAATTCTGAAAGCATGCCTCTTTTGTAATTATGAAATACAGAAATATTTGAAATTCCCCAGTCATCTTGG
CTCACAAATAGAAACATTTGGTTGCGTTTGGTTCACAAGCACACAGGGGCCTGCTGCTTTTGGTTCTGTGTAAAGCCCA
TACTATTTCTTGCTTTTTTTTTTCCACAATGAGTATGTGGGTGCTAGAATGTTATCGAAATATTTTGCAAGACATGAATC
TAAAATGAGCACTCTGCTATGGTTGCAGTTACAGTGCAAACCCCATCAAGTTGCTAGCATTGGACTGATAGACAAGGTA
GGGGCAGCCAGAGAGTCTGAGTTCCTGTGAGGAACTGCAAGGGGGAAGGGGCTGTGTTCAGGGGCTGTCCTGCAGGCTG
AAGGTTGTTATTAGTTGTGTGGTTTCTGGGCCCACACAAAACTCCACCTTCTCCCATTAGAAGTCAGTGTCCTGTTGAT
GTGACCACCTGCTTTATTATTCGGGGTTTTCTGCCATCTAAGACCTGGGCCACTCACCTGGATTTTTGAACTCTTGCTT
TTGCACTGATACCTGATTCAGAATCCGATCAATTGCCTTTCCATCTTTGTCTGTGGGGACCAGAAGTACAGAGGTCTTA
TAGGTGAGCTTTTATTCTTTCTTCCAGAAAGAGTATCCTATAACTGTGTCAGTGAGGAAGTGAAACACAACCTCCTGTT
TGAAACACAAAAGGAATTTAGAGAGGAGTAAAATACAGGGCCGTTTAATTATTCCCTTTAATAAGAAGCAGCATCCCCA
CCAAAGAACATGTCGATCTTACTTGGTGATACATTTCCCTATGGAGAAGGGAGATCTGATTGAAAGAAGACCGAAGGGC
AGGAGGGAGGGAGGGTTGTAAAGCAAGGATGTGGCTCTGTGTAACCGGGATAAAAGCACCCTGCCTTAGGGAGATGTTG
TGAGAAATACCCCAGCACCCTCCTGCTACAAGAGGCAATCCTTATAGCTGGCTCTTGCCATGGTTAAAATGATTGGGAC
AAAAAATATTAAAAAAGAAATGCTCCTCGCTATACTGGTTGGAATCCTCAGGCTTGGAGTAGGAGGAGCTGAGTAATCC
CTGCAGTGTCCCGAAAGCTGGAATCCTGAGTAGATTCAGGCCAATTTGTGTGATATTCCCAGGGATTCCCTCCCTATTT
TGGGCAGAGGAATGGATTTGGATGGGGCAGCATGTTCCTGTGAGTTGGCAGGCACCCACTGAACAGAATAACTGATTTG
GTACCCAGACTGCAGATACCAGCTGAATTGCCAAGCAGATATTGCCTGAGGGAAATACCTGCTAAAGTTGAAATGATCA
CTACTTGGATGGGTTAAACATTTACCAGTATCAACTAGTATTATACATAGTGTTTTAGTTTTTCTGAAAATGTGCATGC
ACATGAGAAGAATAAGTATTAAGATGAAACTCCCAGTCCCTCTCCAGAGGCACAACCATGGTTACCAATTTCTTATATA
AAGTATATATATAAAAGCTTATATAACCTCTTCTTTTACTTATTAAATACTTGCATTGATTACTTTAGAATGGGCAAGT
AAAAACTAGTTTAATTGTAATAATTGTAATCACTCTAATACATTTTCTTATTCCTGAATATTTGCTTTGTTTTAAACTT
TAGCAATTATGAACCTTATACTAGTCTAGGAAAATTAATCTGTAAATATTTCTGTTGGAATCTCAGATTTATAATTAAA
AAATTATATAGGGTATGTTACATAATTTCTGATCAAAGATTTAGATTAAACAAATATGTTTGTGCACTTATTACATCGT
CAGCACTGTCCTGTGCCTCACTCAATACGTGAACTTATTTAATGTAAGTCCACTTAATTTCACTGTAGAAACCAAGGAG
AAAAAGGTCTGGGGCACATACATGGCATTCTTGTGAATCTTTCTGAAAGTGCTCTCTTACCTGACATGGTAGGGATATC
CTGAAAGGTAAGAAGGGAAACCTTCAGGGCAGTGCCATTTGCAGCTAGGTCTTTGAGAATAAGAATGAATTCTTATTCA
ATATGCTGGTGCCACAGTTAGGGCTAAAGTATCTTTAGACCTGCCATCTCTAATAATAACGGCCATTTCTGACTTGTTA
ATCTTCCCGGCTTTTTTCCCTACAATTTTAGGTCTAAATCTTCCATGTAAACACACTCTTTCATGCTGAATCCTGCTCA
AATAACTGCCAAGATGTTTGTTTAGTAATCCTGAAATAGAAGATTGGCTTCTATGCTAAGATAATTATTGGGAGAATCT
CTTCTTTTAATAAATACCTGGTTCTTCTCAAACCCATGCCAAGAAGAATTCTTATAATATCATTTCTTCTCAAACCCAT
GCCAAGAATAATTGTCACCATATCATTTCTTGCTATTCATCCCTTTTAAAACTAAGAAACAGGCCAGGTGTGGTGGCTC
ATGCCTGTAATCCCAGCATTTTGGGAGACCGAGGCCGGTGGATCACCTGAGGTCAGGAGTTCAAGACCAGCCTGGCCAA
CATGGTGAAACCCGTCTCTCCTAAAATACAAAAATTAGCCGGGCATGGTGGCGCATGCCTGTAGTCCCAGCTACTTGGG
AGGCTGAGGCAGGAGAGTTGTTTGAACCTGGGAGGCAGAGGTTGCAGTGAGCTGAGATCGTGCCATCGCACTCCAGCCT
GGGCAACAAGAGCAAAACTCTGTCTCAAAAACAAAACAAAACAAAAAACTAGGAAACATTAAAGCTTCCAAGGAATTG
AGCGTTAAGTGGGGTCTAACATTTAATATGACTGTAGAGTAACTCCTCTTTACTAAGTCTTTTTACTAGTTTCTCCCTT
TCTCTCAAATATGTACAGTTTAAATTGGATGGGTAAAGCAAATGGGAGTATTGCTACTTTTTAGTCATATTTATACA
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TACAAGCTTACTGAAGAATATAACTGAGGCATTAGGAGTGCTTTTCTGATCGGATTAATGTTCAGTTTTTATTTTCAATT
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GATGATGAAACATTAATGGCTCCCTTCCTCCAAGTAGAAGATAAGAGTAATATACGGTCATGGAGTTCCTTTTTAAGGG
GATGTATTAGTTTTCTATTTCTGCATATAAAGTCACCCCAAAGCTTTGTGGCTTAAAACAATAATATTTTATTCTCTCA
TGGTTTCTGTGAGTCAAGAAATTGAACAAGGTAGGGTAGAGATAGCTGGTCTCTTGTTATGTAATGTCTAAGGCCTTAG
CTGGAAGACTCTAAAGGTTGAGGGCTGAAATCATCTGAAAAGTTGTTCATGTCTGGTGGTTGATGCTGGCTGTGGGCTA
GAGACTTCAGTTCATCTCCATGAGTTCTCTCCTTGAGGGTTAGTTTGAACTTCCTCATAGCATAATAGCTGATATCCAA
GGGCAAGCATCCCCAGAGAGAGTAAGAACCTGTTAGAATCTGTGTCCTTTTTATGCTTTGGCTTGGAAACCACAGAGCA
```

Fig. 6.143

TCACTTCAACCACACTTCATGGGTTGGATCAACAACAAGTATCCTTCAATATTCAAAGGAAAAGAACGTAAATCCAACT
CATAGATTCTACCAGTCACAACGTAAAGTAACATAAGAAGATCTGTATATTAATCTTTGGAAAATATAATATGCTATGG
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GAGGCCTCAGGAAACTTACAATCATGGAGGAAGGGGAAGAGGCATGTCTTACATGGTGGCAGGTGAGAGGAAAAAGAGT
GAAGAGTGAAGGGGGAAGAGCCCCTTAGAAAACTATCAGATCTCTTTAGAATTCACTCACTATCACAAGAGCCGGATGG
GGGAAACTGCCCCCATGATCCAGTTACTTCTACCTGGTCTTTCCCTTGACATGAAGGGGTTATGGGGATTATAATTTAA
GATGAGATTTGGGTAGGGGCACAAAGCCTAATCATATCAGGGACAGAGAAAGGAGAAGTGAGGAGGCTGACATTATGAG
ATGGATTTTGAGGGAATAACAGAGATGTTTAACGGGGATATTTATGCAGCTTTGCTGTGTGGAATGTGAGGTAAAATTG
TTTCTCTATATTTTTATAATCTTTAGAGGAAACATCCCACTGGGTTGTAGAGTTGGATCTATTTTTGGACAAAATACAT
TAAGTTGCTGAATATGGCTTCTATGTAGAGGGAGGGAAAAAGTAGAGGCCTACTAAATCTAGGTTACATCTGTTACATC
TGCTTTCAGAAAAACTTAGTAATAAATGTGGCTGTCATGTTAACGTTTGGTGGTTCCTAAATGTCATTGCAAAGACTGA
AGCCAGTCCATGGTAAAATTTTCATTGGCAAGCTCTGAAATCAGAATTAGACATCATGAATCTTTCACAAAGCTAAAT
TTGTTCAATTAAATGAAATGATCTTTATTGTGAGTTTATATCCTTCCAAACTTTTTAGTTTTAAAATGATCTTTTTTAA
GAGGAAATGATGGTGACAGTAGATAGTACTTGTGTTTTGGATTTTTAACTGTCCATATTTGGAAAAGTAAAAAGTTACT
GACCACCATACTTTAGTTTCTAAAACACTGGAGTGCAATCTCTAAATTGCAATGTGTTTGGGGAAAAAAATTCAATCCC
TTTGCATTAGGTATTTTTATTGACATTAGATATAAGAGAATAGTATAAATTTACCTAAATACACAGAAAAATAATCTTT
ACTCTTTATAATCCCAAAGAAATATTCCTTAGTTCTTTTTTCTTCTATTTACTATCAATATCTATCACAAAGCTGATA
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TTTAACTACAAAAGTCTTTTTAGAGAAGAAATATTCTTATCCACCTCATAGCCTAGACTCAGTCAAGGGAAAGGGCAGC
CAAGAATATCTTAAGTGATATCCTGGGCTTTGTTTTCTTGATTCAAGTAGTGTTCACATTATTATAATTATATATTCTT
TTATCTGTATATCTGTCTATTCATCCATTTATTCATTTGACAAAAATTGAGTATCTGTGTTTGAGAGACACAAAGATAA
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TTATCATGTCTCATAGTTAAACCAGATAGTTTGGTAGATTCTTTATCACGTGTGTTTCTAATACAACCAGGAAAACCTA
TCGTATATTCTTTTTTTTTTTTTTTTTTGTGAGACGGAGTCTTGCTTTGTCGCCCAGGCTGGAGTGCGGTGGCACG
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GCTTCCGCCACCATGCCCGGCTAATTATTTTTGTATTTTTAGTAGCGACGGGGTTTCACCGTGATAGCCAGGATGGTCT
CGATTTCCTGACCTCGTGATCCACCCGCCTTGGCCTCCCAAAGTGCTGGGATTACAGGCATGAGCCACCGCGCCCAGCC
TATCATATATTCTTACATTCTCTCTCACACAAATGTGCATGCGCACACACACATGCACACATATACATCATTCTGAC
ACTGCCTTAACTTTCAGAATACAAGGTTGAGTCCTCTGCCACATACTAGCAGTATAATCCTGGATGAGTTACTTAAACT
TACTTTGCTTCAGTTTGAGGATGAAATGAGAGAATATGTGTAAGACATCTGGCACACATAGTGAGCATTCAATAAATGT
TAACTATAAGTAGGTGACCAGTTGGGGCCCAAGGAGATTGTAACTTGCTTAAGGTCATTAAGGTAGTTCGTGGCTAAGT
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GTGTTAACTTTCTGTACATGAAGGGTTTGCCTTCCCAAGTAAGTTGTAAGCATCTCAAAGGTTAAAAGCATTGTGAGTG
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TCCTCAGAGAAAATCATACAATTTTCTACTATATTTCTGCAGTATGAAGAAAATAATTTAATATACCTATGCTTTGGTT
TCTCCATATACAGAACTTATATATGGCATTACTCAATCACCAAAGAATTCCATTATAATTTAGCTTATTGTTATGTAAA
TTCAGGCATAACCCAGGCCACAATATGATCCCAAGAATATAACAAGTATCCATAGGGAAAACCTGGTAGAATAACTCAA
GAGTGGGAATTATTTTAGCTTTGGATGACTTTTATATAAAGGAGCTCCACTCACAATACATTAGTGGCACTATTAACT
ACACTCGCTAGTAAGCTAATATTTGAGGTCCATTTTTTACATCTCTTTGAGAGTATATCAGTATATCAAGCATTACAAA
TTACTACCTTGATTGCCTTTGGAGCTACTCTTTTTTTGAGACAGAGTCTTGCTCTGTCACCCAGGCTGGAGTGCAGTGG
CACCATCTCAGCTCACTGCAACCTCCACCTCTTGGGTTCAAGTGATTCTCCTGCCTCAGCCTCCTGAGTAGCTGGGATT

Fig. 6.144

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ATAGGCGTGCACCACCATGCTCAGCTAATTTTTGTATTTTTAGTAGAGACAGGGTGTCACTATGTTGGTCAGGCTGTTC
TTGAACTTCTGACCTTGTGATCCACCTGCCTCGGCCTCCCAAAGTGCTGGGATTACAGGCATGGGCCACCGTGCCTGGC
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TTTCAGTTTTCTCTTGATGCTCAGGATGAACTAAACATCTAGACAAGGGAGCATCTGGGAAGTCTGACTAACTGGACTA
CTTGGACTGGTTTGTCCTCACAAAATGTAAGACTTACAATCTGTAGGCTGTTACCTAGCTCCTGAGAGCTACCTCCCCC
GGGCCGCCCACACTAGATTTTTAGTCCTGAGCTGCATACCCAGGAATTCTCCTGTTTCCGTGAATCAGGAAGGATATGG
CAAGTTAAGAGAGAAGGAAAGAACCTGCTGAGCCTTGCCTTCCAAGGGATATCAGGAAGTTACTCAAAAGACGGCAAAG
TGCTCTCTACACAAAGTGCTGGAAGAACTGCAGTTATATTCCCTTTAGGAAAGGAAACTTTAAGATGATTTGGAAATA
GCCAAATGTAATCACATAACAACCAATTTTGAAGTTTGCTGGCATGCAACACTCCTTTAGGAATTCTTGGTTTCCAGCT
TGGTCTCATTGGCAACAAATAGAATTTCCTAAGCCTTATTAGGAGCCTGACATTTTGTTTGAGAATCAAAACTTTTCTG
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CGGTTATACAAGGTAAAAATGAGTTACTGGCCCTTCAGGATGGGTATCATGATAGCTCATTAATTTAAATAACTCCTAA
TATGCGTGTAGCAGTTTATCAGTTAGAGAGTGCTTTACATGGGTTGCTTTTCACAATAGATAACTTAGGGCAAGTGAAG
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AGCAAATGGCTTCTATAGTTGAATAATCTTTCAATAAAGTTAGAAAAATGGGTGTCCTTCTTCAATGCATAGCATGTTG
GGAACCCAGGATACTTTAGAAAGTCTCTTGGTCCAGCACTCCGACTTGTGAAAGACAAATATTGTCATACAATGCTTTG
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CATCATTAAGTAAAGGGTGAATTGCACTGGGTTGTTTAAAAGCATGCCCAGATACTGCTGGGTGCGGTGGCTCACGCCT
GTAATCCCAGCACTTTGGGAGGCCAAGGTGGGCAGATCATGAGGTCAGGAGATCGAGACCATTGTGGCTAACACGGTGA
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GTTCATGAAATGTATTCAAAGAAATTTTAAAAATAGCTCAGGTACAGAATCTGTTAAAGCAACAAAGCAGACACATGAC
TCTTCTTAATCTTCCACACTTGACTCCACGGCATTAAGTTAACACTTCCTTAGTCTCCTCTGAGTTGAGACAAATTTCC
TGAAAGGAAGGTTGACATTCCCTGGACACTTTAAGGATCTGTTTGTGCCTCATGGGTATAGGTTGTAAAAGTTCTTTGC
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TTCTTGCTTCCCTTCGTCTGCCATAATAGTCCCTGGGGTATAGTATAAGTGCTTCTAGGGCACTCCTCTAACTGCACGG
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GCAGGGAACAATTGGAACACCTAAAAAAGGGTCAGAGTCCAGAACCAGATTCAGGAGGAGTTCAGCTTTCAATTGGGT
TGGTCCATTTTTCAGTGCTGCCCGCACGAGCTGTCAGTTGCTGGAACTGCAGTATTTCAAGGAAAGGCGCGGCTAAG
CAGAGGTGAAATCTTTAGACAGATGAGTCTCAAAGAGGAAATTATGCAGAAGCCAAATGGCTACAAATGGTACTGAGGA
TTGAAACCTAAACTATAAGTGAAGGTCTCTTCCCTAAATGATACACTCAGACTTAACAATAAGCTAATAAGGTGGTTCA
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TGTTTTTGGTATTGATGCCATTGTGTTGGGAGTACAGCAGAAGGACAAAGAACTCCTTTAGCAACTTGGCCTGCAGTCA
AATAGGCCAACATTGGAATCTGAGCTCCACTGCCTTCTTCACCTGGGACTAATCACTTACCCTCTCAGTCTCCTCATCT
GTAAAAAAAAAAGGATAATAATAATAATAATTGTTACCTTTGGAAAATTACACCTTTCTCTTCTGTACCTTTAACCTTT
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Fig. 6.145

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CTCTCAACTAGATACTTTCCATGAGACTCAAAATACATTCAAGTGTCTTCTGCATTAAAAACAACAAAACAAAAACCT
CACCTTCAACCACTTATTTCCCTATGTTAGGGTTTCTCTTGACGTTTGGGGCTAGATAATTCATTACTTTGGGGCTGTT
CTGTGCGTTTTGGGATATTTAGCAGCATTCTTTACCCATGAAAATATTGTGTCCCTCGCCCCACAAGTTGTGGCAATCA
AAGTTGTCTCTAGACATTGCCAAATATCTCCTGGAAAGATGCCCACCCCCCACCCCAGTTAGGAACCACTATTCTAATG
ACTGCCTCATCTCTTGTCTCCCTTGCAAAGCCACACTTGCCCAGTTTTGCCTATTCTCTCTAGCTATATTCCTCACTTC
CCTATCACAGCCGATCCTATTCTAGTCTGTCTCTTGTTCCCATTTTACCAATAAAATCATTTTTAGTAAGAAGAACAAT
AACCTGGATAGTTCTAAATGTAATGAATATTGCAATTCTTCTCTTCCTTGATTTCTCACAATTTGAAGTGTGTTCTTCT
CCTGGCTCCTCTGACACTATTCTCTCTCGGCTTGATCTGTGAAGCCTGGGTCTCCTGGAATAGCTGCAGCCTTTCTCTC
CACTTTCTTACCATATTCTCTCACTGTGTCCAGGACACTAGCTCAGACTTTTTCAAATGCTGATTAAAGGGCTTCCATG
CTGACAGAGGGCAAACCTCACAGAGCAAGCTTTTCTTAAGCTTTTGCTTATATCACATTTGCTAGCACTCTCTTGGCCA
ATGCCCAGAGTCACATGGGAAGAGCCTACTGAAGGCAGAAATACAGAGAGAGAAGACTTTTGGCATAACATCTTGAGGG
CTTCCTTTAACCCACTCCCCAGTAAAACTAGTGAAAACTATGATCAAACAAATAAAAACTCCAAGCACTTAAATCCTCT
GGAAATGATCCTAAGGACAAATAGCAAATGAAGAAACATCTATTTAAGAACATTTATGAAAATTCAATAAGAAAGGCAA
GCCTGTGGTATTTAAACCAAGACTGCTCCCTCTCACCCCCTTCCAAGTTCAGGGAGATGGAGCTTCCATTCCAGGCTGG
TGCAATCAAGAACACAGAGCTCTCTCCCCTCCAGCTGGAAGGTTTCTTCCTGGAAGGAACAGAACTTCAGTGTTTCTCA
TCCTGGCCATAGTTACCCATTGCTAAGGCTAAGCTCTGGTGAATACAGTAGAGAGGTAGGGGCTTCCTCCCCTGCCAAA
TCCCCCATCATTGAATGGAGGGGATACCTTAGGCACTGCATGCTAAGAATACAGAGGCCTCATCATCCTTGCCTGGCCT
CCTGAGGTGGGGGTTCCACACCAGGAGAGATAAATATAGAAGATATTAGAGTGCTGCCACCTCCCAACTAAGCTAAGCT
CCTAGAGTGGGAGTTTCATGCAGTCATGCAGGAAGAACCTCTCCATTTTCTCCACCTCCATCTTGAGAAACATGGCTTA
GAGATTTTGCCTGGTGGTGGCGGGGGCAATAATCCATCATAAAATAAATATTCTTAATCTCTTCCCAAAGGGCATGGCT
TCATTTGCAACAAAGTATGGAAAAGTTCAAGCCTTAGAGTGCGCTCAAGAACAGTGGAAGCTATGGTGAAAGGCAATTG
GGAGGAGAGTCAAGATACAGGCTAAATTGCAGACTAGTTTGCAGGAGAGAACCAGAGAAATAACACAGCTGGGAGGAGC
CCTGACTTCGAACTATTTCTTCAAAGGAGACACAATTCGATTGTATTAGTGTGTTGAACAATATAAGGTTGTAAAGCAC
TGTTGAAAATAACACAGCAATTGTTCACCAATTAGTGGAGTTGAACAGCTGAATGTGGTTAAGGAAAGAGTGAAGGACA
ATCACACGAACATCACTCTCATGCCGGCTGGGCTGTCTGTGGACATACCCAAAGATGTACCTCCCTGATAGCAAACTCA
GTGACTTAACTGAGGGAAATTAACTCACCTGACTGTGAAGGAAATATACTTATTAAAATAATCCAGCCCCTCACTAAAC
AACCAAAGAATAATAGCAAGCCCTGGGTGGTGGTAGGGGGAGTAAGAAGAGTTGCTACAGTATATTATCTGCAATATCC
ACTTTCCAACCAAAGATCACAAAGCATGGAAAGAAACAAAAAATATAACCCATACACAGGATAAAAGAACCAGATGGCA
GAAACTATGGGTGAGAGGGACCCAGATGTTGGATTTAATAGAGAGATGGAGAGGAGAATGAATGCAGCCAGTGTGCAAT
GTGTCATGTCCTCTAAATGTTCAAGTCCCTCAAGTCATAGATCCACCTATTTTTTCTCAAGTTAATATCTCTACACCTA
GTTTTAATTCCCTGTCATGTATACAGGACTCACACATTTTATCTTCAACTTAGGCCTTCCCTCTGAGCTCCGGACCTGT
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CTCTAACAGCTCCCTGCTTATCACCCCACCCCATCATATTGAAAACCAAACCCATTTAAATACCTTTCGTTTGTGTCCT
CCTCTCTCTACCTCCACCACCAGTACTGTAGTGTAATCTGCCATCATCTCTCACATGGATTACTGCACAAGTCTCCTAT
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ATTCAAATCTAATCATGGTATGTGCCTCCCCCTTCTTCTATGGAGACCATAGGAAAGAAACCGAAACTCCTGTGGTGCT
TGAGTTGATGCTAGTAACCTCTAAACCTCATTTCATGCCTGGATCTGTCACTTCCTGAAGCTCCATGATAAAGCACAGC
CTCCTCACCCTCCAAGTGCGGGGGTTCTCTAGTTGTCTCTTCCAGGTGCCACCAGAGGGACTTCTCTCATAAAAGGCCT
GTGCACCCTATGCTCTTTCTCTCTCTGTTATGGACTTCCTTCCAGAGAGCTGGTCCCCATTCCTTGATATGGGACTGTC
CACTTGTATGAGTAATAAAAATCCTTCTCTGATCATACCAAAGAATTGGTGTCATGAGTTTTTATATTAGAAGCTAAAC
TTGAGAGGGAACTTCCCCATGTCACAGTGAAGGTGTGATGATACCCTCTTTCTACCATCTTTAAATTCTTCCGTCGCTT
CCATTCCATTTTCTCTTAGGATAAAGTTCACAATCCTTAGGTTGGCCTACAAGGTCCTCTACAGTTAGGTACCCCGTC
TGCTTTCCTTCCTTCCTTCACATTAGTTATGCTCCCCACTGCCTGAAAACATGCAGACCTGCTGCATCCTCTACCTGAA
ATACTCTTCTCCTCACCCCTGCCAGACTTTACCTAATTTGCCCTTCGTTTGTCTTTGTAACACCGCTCAGAAATCTCTT
CCTCAGAAAGGCCTTCTCTGACACCTCAGCACAGTCCTCCTCACGAGGTCAGATGTATTTTTTCTTCAAGGCACTGACT
TTGGTTTAAAATTTGTATTTATTCATGTAATTATTTGGAGTGCATTTATCTTCCTCACTAGATATAATCACCATTAGCT
CAAGGGCTTTATATGTTTTGCTTGCCATTGTGTCCTCAGTGCCTAGGCCATTCACTGACCCACAGTAGGTGCTCTGATA
AATATTTGTTGAGTGTAGAATAAATGAAGACCCTTAGAAGCATAAAATATATAATCTCAAAGCAGAAGATACTTTAGTT
AGAATAAGTTAGGTTCAACTATATTAACACTCCTTACTACACACACACATAGCTACACTCATCAATTTTGGTGGCTTAC
ACCAAAAAGGCTTATTTCTTGTGCACACTGCATTCTTGTGCTTTAACAGGATACTCCGCCCTACCTCATCTTTATTACA
GACCTTGGCTGACGGAGGCTACCACTTGGAATATTGTCTATTTTCAAAGTCCAGCAGAAGAGAAGTTAAGAAGGGTTTC
ACTCTGGCAATTAAATATTCCAGCTAGGAAATGACATACTTTATCTTCATATACAATCCATTGGCCATAACTAGTCTCA
TGGTCCTGCCCATTTGTAAAGGGGAAATAGGTAATCATCTCATGCATTGAGGAGGAAAGAAGAATTAGATACATGTGAC
CAGTAGAAGTCTCCTTACTGACATTTTAATGTAGAGCTGTATGTGATTAAAGTTAAAATTGAATCTGAAGGGCTGACAG
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TGCTAGCAGGAAAGACAAAATAAACATAATCAAGAAGTATGTATGAACGTGTGCCCAGGGCAGTGAGGAGTGATCAGCC
TCCTTGAAGTGGAGGATGTGTGTCAAGGAACCAGAGGAGATGATTAGAGTGGCAGACCTGATTATTAGCATATCCATGC
TTCTGTCTGAAGCCTTTTTGGCCTATAATGTAGAAATATCCTTGGAGTAAGCAGTTCAGCATATGGTTTTAGACATATA
GGTCATCTTGGCTGGATTGCATATTCTTCATTTTTAAAATTCCAGGCTGTATTGGTTGATATGCAGTCCAGAAAGAGCT
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Fig. 6.146.

GGATCTGACTTAAGAAAATAAAAACTGCTTTAATCCCTCTTTTCAGCCCATTGTCTATCCATTAGCAGGTAAACACCAG
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CAGCCTGCCATGATTCAAATTGAAACTCCATCATTCACCACTGTGTGACCTTGAGAAAGTTATGTAATCATTCTTGGAT
AGTGATTCTTACTGCTATGGACAGATTTGTGTTCCCCCCAAATTCAAATGTTGAGGCAACTGTGTGAGGAGAGAGGGCC
TTTAGGAGGTAAAGTTAAATGAAGTCATAAATGTAGGGTCTTAATTCAATAGGATTAGTGGCCTTATAAGAAGGGGAAG
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CAGGGATAATAACAGTGCCTATATGATAGTGTGTTTACAAGGATTTAATATATGTTAGTTTATATTATTTTATGTCCAT
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GACAGACCTGGTTTGGATTTCAGTTCCACCTCTTGTAAACTCTATTACCTTGAGCCAGTGACTAATGTATCTAAGCCTC
CATTTTCCTATCTGTAAATGGGGATGATAACTAGTGCTGCTTGTCTCTTCAGGTTGTTGTGAGGATTAAAGGAGATATG
CATGACAATTCATCTGCCAGGTAGTAAGCATTCCAAATATGCTATTTACTGCCATCATTAGAGGTTTGCTGAGCTTCCT
CTTTTTGCATTAAGTAAGAGACTATTCTCTCCAGAAAACTTTGAACTACATGATGGAGGAACAAATAATAGCAGTTTCTC
CTCAACCTTGGAACTCTAAAAATGTTTTTTCTAAGCCATTCTTATCTTTATTTTGTCATTAAAAGATACATGCATTGTG
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CAGTTATTATCTCACAATGTGAAGGCAGGAATCTATAGACAGATTATTGAACATCTCATGTATCATATCATGTATATCA
TTAAGCTTATATGTATGACAAAATAGTATTTTGTGCAGCAGTGCTTCTCAAACATTAATAAATGAATCACCTGAGGATC
ACATTAAAAATTCAGATGCTGATCAGGGAGTCTAGAATGGGACTGAAGTGCTGCATTTCTTTTCTTTTTTTTTAGACTTT
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GCAGGAATCAACTGGAATTAGAGTGATAGAACCACAGCCCTTAGACATGTATCAGACACTCCCTGACCAAGTCTGTCCA
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CAAGGTGGGCGGATCACCTGAGTCAAGAGTTTGAGACCAGCCTGGCCAACATGGTAAAACCCTGTCTCTACTAAAAAT
ACAAAAATTAGCCAGGTATGGTGGCCAGTGCCTGGAGGCTGGGGCAGGAGAATCGTTTGAGCCTGGGAGGCGGAAGTTG
CAGTGAGCTGAGATGGTACCACTGCACTCCAGCCTGGAGGATAGATTGAGACTCTGTCTCAAAAAAAATAAAAAGAAAA
AGAAAACCCTCACCTTCTATAATCAAATAGGTTTTTAAGTGTTTTCCTAAGCTTGATAGCACTTCAGTCTCTCAGAGGA
ATTCTGCAGTCCTGCAGTTGTCTTGGGTCAACCAGCCAAAAGTTAACTAATTTGCATAATGGCAGCCCCATAAAATGCG
CAAAATTATTATCTTTTATATGAATGCTTATTAATATATATATATATATATATCTCCTTTTACATCTATCTAGAAAACG
TTTTAAGGGAAGAATCATATAGCTTCAAACATTTTTGTTAATTACAATATGGAAGAAAAGAGAGTTCAATTCCTTAATC

Fig. 6.142

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TATCATGTGTGAGTTAGGGGACTTTAAGCAAGTTATTTTCTTAGGAATCCTGTTTCCTGACCTATCAAATGAATGTAAG
AATAGCTACCTCACATAAGTGTGAGGACATAGATAATTTCTATAAAATCATTTAGTTAGAGTCTTGGCACATAGTAGGA
GCCCTGCTCTCAGTACAAGTTTAAAATGTCTGTTGAATGCTAAAAGATTACTCCATTGAGCTTTTTCAAACTTTAAAGT
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GACACATTAGTTTAGAGACATATCAATAATTATGGAAGTATCTTCTCAAACTGTAATGTTTTCCTCTGCCTTAGTGCTA
GCAATCAAATCCACTGAATTAATCTGAGCAGGGTCAGGACTTTGAACCTGAGGAAAATCACTTGAATTTGAGGCATTAA
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GGTTACTGATGATGGTGGGGAGGTCTTTACAGTGTGATGGCTTGAAGTACAAAGTACAAACTTGAAAGTCAGATTACCA
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CAAGAGTTGCAAAGTAGAAAACTGTCAGTGAATGTCAGCTCTCTCTATTATCCCTCAGAGACTCCTCCTGCCACTTTAG
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Fig. 6 :148

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GTAGGGATATATATATGATCTAATATATAGAAATGGATATATAGAATATATATGTATATAGAATATATAGAAATTTCTA
ATATATAGAAATTTCTAATATATAGAAATGGAGCATTTTAAGGTCTGGAACTTTGGGGCTGTCAACAGTTATAAGAAAA
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GCGCGCACCACCACACCTGGCTAATTTTTGTATTTTCGGTAGAGACGGTGTCTCAGGATGTTGGCCAGGCTGGTCTCAA
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TCAAAACTTGTTGAAACTCCCTCTCCATCAGATTCTGTGTGCCCAAAGAACTGTCCTTGGAGGAGAGTTTCAAATGTCT
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TTCGTACTAATTAGCACCTGCACTGAATTCAGAAAGCAGCACTTGAATCAAACCAAGTCAAGCTGTCTGTGAAGGAAAA
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CTCCCATACCTCATAGAATGTGTTTGGAAGTATTCTCTCCTCTTCTATTTTTCAGAATAAATTGGGAAAATTGGTATTA
CTTCTTTAAATGTTTGGTAAAATGCAGCAGTGAGGCCATCAGATCCTGGGCTTTTCTTTGCTGGGAGACTGTTTATTAT
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TTGGGAAGTTCTCTGTTATTATCCCTTTGAATACACTTTTTATCCCTATGTCTCTCTCCACCTTCTCTTTAGGGCCAAT
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<p align="center">Fig. 6.149</p>

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TCTTCTTTTACTTTCTACCAAACAAATGGGGTCTGTCTCTCTCTCTGTCTTGAGCTACCTGGAGCTTGGGGAGGGGT
GACACTAGCACCCTTGTGGCCACCACCACTGGGATGGCGCTGGGTCAGACCTGAAGCCAACACAGCATTGGATCTCTCT
CAAGGCCTGCAATGACCAGTACCTGGCTACTGCATATGTTTGCTCAAGGTCCTAAGGCTCTACAATCAGCAGGTAGCAT
AGCCAACAAGGCTTGTATCTGTCCCTTCAGGACATCAAGTTCCCCTTTGCCCTAGGAGAATCCAGAGATGCCATATGGG
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GGGTCACACCTGAAGCCAGCATTTCTCAGTCTCACCCCAGGCCCACAGTGAGTACCACCTGGCTATTGCTGCTGATTAT
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GGGAAGGAGTCTCACCTGCAGCTGCAAGCTGCTCTGCCTAGGGTTGGGGGTGGGGTGGCACAAGCACTCCTTTGGCTGC
CCCTGCTAGTGTCTCACTAGGTCACATGTCCCCCATAAACACTAATTCTAAACCCATCCCAGCATCAGTACTTGCCCAG
GAATTTCAGTCTTTGTGGCTTAGACTGTCGTTCAAGTTTATTTAGTACCCCAGAACACTTTAACTTATGGCAGTGAGGC
TTGCTGGATTTCAGATTCTGACTGCTGGAATGGGCAATTTCACTCTGGCTAGGGCTGGTCTATATTCTCCCTCCACTGG
CGCTAGCTGAGTTCTGCCCAGTGTTGCTTTCCACTGTGACAGGGCAGCACTGAGTTCCAAAGCAAAGTCCCCCAGTCGC
TGCACTCTCCCTYCCCCGAGCACAAAGATTCACTCTCTGTGCCACGTGGCCACTGCTGACGCATGGGGGAGGGGTGGCA
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TCTTTTGTTGAAATTCTCAATAACCTCAACTTTTGATTTCAAATGTTAGGCCTTATTATATTTAATGAGAATTAACACT
ACTTAATAATGTTCATATTATATGAATGGCCATTATGCTGTTTTCGTGTTTTAAAAAAAGTTTGCAAAAAACATTAATT
TTTTGCCTAAATTATTTTGAACATTAGACATGTTATATTTTAGCAACCCTAATAATAATAATTTTTAATTATTACACTT
AATATCCTGCCCACTTTTGAAATATTGACATCATAGAGCTTATGTATGAGACAGGTAATTAAGTAGTTAACATTTAGTAA
GGATGTTTTATGTGGCAGGTACTATTCTTAGCATTATACACCCATTACCGTACTTAATGTTCACACCAACACTATGAGG
TAGACACTTTTTTTAATATACTTTAAGTTCTAGGGTACATGTGCACACGTGCAGGTTTGTTACATAGGTATACATTGC
CATGTTGGTTTACTGCACCCATTAACTCGTCATTTACATTAGGTATATCTCCTAATGCTTTCCCTCCCCCTCCCCCCTC
CCCCCACCACATGACAGGCCCCAGTGTGTGATGTTCCCCTTCCTGTGTCCAAGTGTTCTCATTGTTCAGTTCCCACCTA
TGAGTGAGAACATGCAGTGTTTGGTTTTCTGTCCTTGCTATACTTTGCTCAGAATGATGGTTTCCAGTTTCATCCGTGT
CCCTACAAAGTTATTATCCTTTCCAGATGAAGAAACTGAAGTTCAGAGAAGTTAAGTGGCTTACCAAAAGTCACACGAC
CAGGATTCAAGCCTGGATAATGTGGCTCCAGGGTCTGTTCCCCTAACCACTATTTCATAATAAGGACTATCTTCCAACT
AACTCTGTATGTTACATACTGCTAGTTCAAAGCCAGGTAGACGAAAAAATACTCCCTGTTTCAGCTGAAGAAGGTTTTC
CAATATATTTTTAGCTGCAAATATAAAATACCACAAAGCCAATTTACATTAAAGGAGAATTGTACATTCAACACATCAG
CATTTCCACAGGATTGTAGAATAATATATTAGAACAAAAATAGTGTTTGTAATAGAAATACAGTTATTTTACTTTGGAG
AATGGGCTTGGAAATGGCAGAGATAAGAATTATAAGTTATTTATAATTAGCTTAATAATTTCGGTATTCTTATCTGCAG
CCATACAGAGTTATGTGAGTTGTGAACTGGGATAGGACATTAAAGCTGACAGGGTTACATCTGGATCAGAAACAAGACA
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CTCATTTAATAATAGGGCTATGTTATTATAACAAAGCATCTGCTCTGTCTTTCTCTCAACAGACCAAATTAATCTTTGC
CACATTATTCCTTGGACATTTGAGAGGTTAAAAATATATGCGCATATACATGTAAAACCAATTTGAAAAATAAAAGTTCC
CAAATGTTTATTCAGTGTCTACTTTGTGGCAGGCACTGACCTGAGGGCCGAAGAAAAATAGGGAGGGTCTTTCCCACAA.
GAAGGAAGAAAAGGAACAGATGGTGCATGATTACCAGGTTCAGTGGTTTGCATGGAGTGCTCTTCAAGCACTGATAAAA
AGTGTCTAAATCAGATTAGATGGAACAATGTCAGATATGGCTTTTAGAGGATGTGAATCTGACCTCAATTTGAGAATGG
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```

Fig. 6.150

CTGGAATTCTTTCATATCTAATGAAAGAATAAACCTCCTAGAAAGTCCATGCAGATTGTATGGTACACATTTTAATTGG
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TACTTTAAGCACACATCTTAAATCCAGGCTAAAACTCTTAGACTGGCATTGGCAATTAGCTATTTCATGGATATAAACT
AAGGAGACTTGACGTTACCAACCTGCTTGGGGAGTATTTTGCTGAGTCTATCCAACTGTTATCAATTCGAGAACAATAC
TTTTTAATTTGAATCAACCTGTTTTTTTAAAAAAGGTTTTGTTTTAACTAAATTCATCTGTTCTCCCCACAAAAAGCCA
GTTTGTTTGTACTTACATATCTCTATGTATACATGTATATATTTTTGAGTCTGCAATATTCTTGTCAGAATAATTTTGC
TTCCTTGAATATTTCAACTATTCAAAAATTTACTGGGTTAAAAACAACACACACTTACTATCTCACAGTTTCCAAGGGTC
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GCAGTTATAGCTTTGTGTTGATCCATAGACTCTCTGGTTTGTTCCCCATTGAGACCAGCTAAGCCATGAGTCACAAGCT
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GTTTTTGTACTCCACTTGAAGCTTTAATAGATTCTCTCCAACTTAATCTGTCTTAAAACAAACTCTGGATTTACAACYT
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CAATTTCTTTTCCTTTCTTTCCTTTTACACACATACACACACACACWYACAYACACACACACACACACACACACA
CACACACCCCTATCAGTAAATTATAACCAAGCCTTAACTATACTTCCCCTTCTCCTACCTACTTCTTCCACGCCGGGCC
AAGCCAGTTATCTTTCTTAATTGTTGAAATAGCTTCCAATCTGCTATATGTATCTCCACACTTGCCATTCTACCGTCTG

Fig. 6.151

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TTTTCAACATGGCAGCCAGAGTATGATCTTCTTAAAATAAAACTCAGACCATGTCACACTTCTCAAAACCTGTCAAAGG
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ACACACACACACACACACACATACATCTTCCTGGCTTTGTCTTTTCTGATCTCTCCCTCAACTTGCTCCAGCCACACAG
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CTCAGGTCTCTGCTCAAAAATCACCTCAACAGAGAGGACTTTCCTGACCACATCTTGTAAAATTGCATACCTCCCTCTC
CCAGGCTCCTTACCTTTCCCAGGCTCCTTAACCACCCCCCATCTTTGTTTTTCTCCACAGAACTTATTATCACCTGATA
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CACCTAGGACATGTAGTAGTTATTCATATGTATTAAATAAATAAGTTATATGAAATATATCATTTAACAGCTGATTTGC
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CTATTCTTAAACCATATTTAAGAAGGTGAAATAGAGATATGATTTGATGGTTTTTATTATTATATGGTTTTCAGTATTG
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CCTTCGAGTTCAGGGTCCAAAAAATATCTCAAGCACCAATCTTAGGTTTTACAATAGTGATGTTATCCCTAGGAGCAAC
TGGGGATTTTTAGAATCTTGTGACCTCTAGATGCATGATTCCTAAATCGCAATTTCTAATCTTGTGGCAAATTTGTTAG
TCCTACAACAGCCAGTCTGGTCCCCAGGCAAGAAAGAGTTTGTTTTGGGAAAGGGCTATTATCATCTTTGTTTCAAAG
TTAAACTATAAACTAAATTCTTCCCAAAGTTAGTTTGGCCTATGCCCAGGAATGAATAAGGACAGCTTGGAGGTTAAAA
GCAAGATGGAATCTGTTAGGTCATATCTCTTTCTCTGTCATAATTTTTTCACTGTTATAACTTTTACAAAGACAGTTTT
ACTCCTATGATAATCCTTCAGTTAGTTCACTGTCTAGTTCCTGGGAGTCTGTTTTGTGGATCCTAACCCTACCCCAAGG
ATTCTACTGCATTGTAAATTCTGACTTTACATTTAAGACATGGAGTTCTTTGGGGGCAGGGAGATAAATATCCTATTTT
AACTATCAAGGGTTAAAGTGCTTTCAAGGATCTAGTGTCTGGTGGTGATACCAGAGTCATACAATTCAAATTACAAGCA
TCTGTTTGTTGTGTTGAGAGCAGGGACTGATAAATGAGGCACTTTTTGATTGTGCAACTGGGTATAAAACAGTCAAAAT
GATGCTTATAACTTCAAGCTTTTCTCCCACTTGAATCAGCCAACAAGATTTTCATACAACATTTAGACCAGAGTAGCTG
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AACATGAGAGAGGTGTAAGCTCATAAAATAAGAAAATCTGTTTTTGTTAAAGTAGACGATGTGCAATTCAGAAACAATG
AACCATATCAACTAAATAATGGGTTAAATCAGTAGAGACTTTGTCAATCTTGTATTGATTTAATCATTTCAAAATTATT
TATTCATTGTCTATAGGTATGCTAGGAGAATAGTTGAATACAAAAAATGTTCTTACACTCAGGAACTTCACAATCAGGT
GCAAAATGAGACAGGCATACTCACATTAAAAATTAAATGACAATTCAAGAGGCGGCACAGGCAGTCTAGGATAAACTG
CCATATGAATAATAGCAACAATAAAGTAGATGAGCTCCCAAAAAGGAAAGATCGAGGAAGGCAGCATGGAAAAATTTGA
CATTGCCTTGTGTTTGGAGTTGAACGAGTTGGAAGTGAATAAGTAGAGGTTTTTCCACGTTGCAGAGAAGTACGATGGT
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Fig. 6.152

GGGGGGTATATATGGGGCATTCAAAATGCATGAAACTAGGAATTCGTTTGTCATCTTTAGGAAATTTCAGAAGTAGTTT
CTGACCATTTTATGTGGGTCCTTTATTTTCAAGTTTAGACATTGGGATTTTATCCTGAGAATAATGGGGAACCATTGAA
GGTTTTCTACAGGGGACAAGGATAGCATGTTGGAATGACTAACCAGATAGAAATATGCACAATTTCTTGAAAATAGGGA
GAACGAGAAAGGAAGAAAGCAGAACAGTGGTTCTTGACTTTAATGTGCCACCTAAGGGCMAGGTGCAGTGGGTCACACC
SGTAATCCCAGCACTTTGGGAGGCCGAGGTGGGTGGATTGCTTGAGTCCAGGTGTTTGAGACAAGGCTGAGCAACATGG
TAAAACTCCATGTCTACAAAAAAATACAAAAGTGGTGGTGTGTGCCTGTAATCCCGCTGCTCGGGATGCTGAAGTGGGA
GGATCGATTGAGGCCACAACATTGAAGCTGCAGTGAGCCATGATTGCGCCAGGGCACTCCAGCCTGGGTAACAGAGCAA
GACCCCTTTGTCTCAAAAGAAAAAAAAAATCCACCTAAGGAGCTTGTTAAAATTGCCGATTCCTGAACCTTACTCCAGA
GATTCTGGTTCAGCAAGTCTGGAGAGGACGCATAATCTCCCTGTTAAGGGTTAGATTCTAAATAAGTGGTCTTCAGGCC
ACATTTTGAATAATACTAAATTAGAAGGCTGTCACAATAATCTAGGAGTGAAATGGTGAAGGTTTGTATTATAATTTTG
GAAGCAGGGATGAAAAGAAAGTGATGGTTGGTAGAGATATAGTGAAGGAAGAATTGATAAGACCTAGAAATGTTTGGGT
ATAAGAAATCAAAGGGAAGAAATAGTTTTAAATGATATTAATGTATTTATCATATTTGATATTTTTGTTAATGTAACTG
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TATAAGATCAAATGATGGGTATGGGAATATCTGTAAATAGGAAAGAAATTCATAAAAAATTAAGCTACCACTTTGTGTT
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AATCATTACCTTTTGAAAGTAGATGTAGAATTCTCAATTATCTAAATCTTCCACATTTAGCTGAGAGACCAACTCTAGG
GAATATATTAGAGTATTTTCATAGTTCCATCAGACTATGAACCCCAATTCTTAGTTAAGGCACTATAGCTATGGGAATT
ATAGTCCATAGTGAATAAAATATACAATCATGGACCTTATTTTGAATAGCTTCCAGAACAAACATATGCGAACATCTTG
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GAATATTGCCTAAAATTGCTGAAATTACCAAGTCTTTAATTTCATCAACAGAAGAAAATAGGCAAAGAAATTCAGGCAA
ATTGAAGAGTTAAAACGTTATGTATAGGTCAGGTTCAGTTCAGTGCAAATAACTGTAATATTCCTTTATGTTTTAAAGG
GATGTTTGGTTTGAGAGGGAATGCTTTCACCTTCTGAAAGGGAAGAGGGAGGTCAAATAGTGACAAAAGAATGAGGACT
TGGGGGAAGTTTCAGCTAAGAGGACAAGCAAAAGAAGGCTGAGAACAAACAGCAATTTGTGGTGGGCATGTTTGGGAGC
TGTAAAGAGAATAGCTCTTGCTTTGGAGTAAACTTCCTCAGTCAGAACTAGGTCCTTGAATTCAGACCAGCTCTCATCT
CCTTGGTCCACTCTAGGGCTGATGGAGGCTGCTGAAATCATCTTTTGCTACATTAATACTCATTTTTCGCTCAAAGATG
ACAAGGAGTATTTAAACGCAGAGAGAATGTTATTTATAAATGATGATTTGTTCGTGATAGAACTTGCCACATAGTATTT
TCTCTTTTTTAACTTTTATTTTAGGTTCAGGGGTACATGTGCAGGTTTGTTATATAGGTAAGCTAGTGTGATGGAGGTT
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AATATTTTCTTATTCATTAGTGGTCAGCAAACTGTTTTTATAAAGATTCGTTTAGTAAATATTTTAGACTTTGTGAGAT
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CTGTATCTGGCAAATTCTTTAACTAAAGAAATGTAAAACGCTTTGGAGGTACCTAAATAGTGGCTAAAAATTGAGTTCC
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AAGAATGAATAATATTTAACTGTGATGAGACTACTGATTTGGTTTAGATTCATCATTTCTAAGTTACTTGACCAACATC
AAAGAAGAACTGGGTCAATTAGTCCAATCTGTGGTTTATAATTGGCCAATTAGTCCACATCTTTGAATCTGCACAGGGA
CCACTTGCACATAAATCTAGATCTACATTTGCCACGCTCCACAAAGCATTCTGAACAATTCACTGTGTTCAAAAATGAG

Fig. 6.153

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ATTTCTATTATTTATGCATGAAGAAAATATGCTGTTTTCCATGCACTGAGCCAGGGGGAAAACTACTCAGCCTTGGTAA
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TTCCTCCAAATACTATGTCAAAACTACTTTGATTAAAATAAATAAAATCTTGTATCACTGTGCCAACTCTGTAAGAAGA
CAGAGCAAGCAAGCAAACACATTAAAAACTAAGGTGTTGGAGAGTCTGCCGTGACTTGTGAATTCTGTATATTTTTTCC
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TCTTTTCCCAATATTTAAAAAAATGTATTATTCATTAAAAATTTTGCATTCACAAAAAAGTGAATAATTGGGCTCCAAT
CTGAGTTAGTTTATGAGTGTTCCCCAGGTACCTATCCTCTCTCTTTTGGAGAGAACATGTATTCCTTGAGGACAGATTA
AGAAACTAGTCTTCCCTTAATGCTGGGAGCCTCTTTGAAGTCTTATTTACTTCAGATCTTAAGGGAGAATTGTGGTACG
CTGAAAAGTATCACATTTGTTATCACTTGAGGAAACACCATTTGATCTAATGAGCTAGACTTTTTCATCTTTCAATTCA
TGACAGCTATAGCTACATTAGAAATTGCATTTTGGAGGTCTTGGATAATTATCTAAAAATATTCACAAACCCCTTGGAG
GCTAATGAATTTAAACTTGGATGGCTAATCCTAAAATGGCTTTATTCCAGCAAAGTGGGAGAGAAACTCCCCTTCTTCT
GTTTATTTAAGGTTTCTAGTTTGGGGTCCTGGTTCTAGTTTGGAGTCCTGGAGTTCCTTTCAATGTGTTTCCATAATAG
CCTGATTCAGAGGAGTCGGGGAGAAAGAAAGAAACAAAAAGAAAATGCCTTGTTTATTTTATTTACATATATACATACA
TATCTATACATATGTAAAATGCACATGGTAAAGCATAGTATTTTGCACAAAATCTTAAGTAAAGACAGACTTGGGTTTA
GATTTATAATAGGATTTAGTAATACTAATACCAATAACAGCAGCAGCAAATTCTTGAGTGCATGAAATGGATTACTTTA
TGTAATTTCTGCAAAGTCCTATCATGTATATTTTGTCACTAGTTTATTTTACAGATGAGGAAACTAAGGCTCAGCAAAG
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Fig. 6.154.

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ATATCATTCTTATGAATATCTTGTTGCTTTTCAATAAGATATGGAAGTGCAGTTTTTCTCCCAGGAAGAAATATTTTTT
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TCAGGTTGGATTTGAATTATTCCAATTTAATCTGGAAAAAAATTATAAGTCACTTGTACTGTTAGTRTATGAGAAATTA
```

Fig. 6.155

161/375

GAATAAGAGTTTCGGATAGAGGTATTTTGCCACAGTTATATTGAGGCGAAAGGTGGCAAGCATTTCAGGGGAAAGAAGA
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GAGAGAGTGGAAGAAAGAAAAATAAGCTAATTGGAATAGAGGTATTGCATCATTGATTTGAATTTATTTTTTGCTATGC
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TGTAGCTGCCACTGTGGCTCACCTGATATATCATTGTCATTCTATTTAAGGATCCTTAGACTTAAAATTCTTTACAATA

Fig. 6.156

CTTGCATATTCCCCACATGCTCACACATTGCATGTCATGTTTTTCCTTAATAGAGTTGCTTTTGCAAATAAATACATCT
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Fig. 6.157

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Fig. 6:158

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Fig. 6 160

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AACCTTTGATTAACTTAAAGTCAACTGATTAAGGATGCTAATTACATTTGAACAATCCTTTAACTTTGCCATATTCCAC
TGGTTAAAAGAAAATTACAGATGTTGACCATCTGGGAGGAGATAGTGTAAGGGCCTGAGCTATTGGAGATCATCTTAGA
ATTCTGCCCACCACAATCGTGTTCTATGTTTATGCAGGTATATGTGTGAGTGTGTATTTATGTGTGTAGTTAATAACTT
AAAAAATGAATTACATGTATCCCAAAACAAGCGGCAACTGCACATCCCATGCCTTCTTACTGGATTAGTTTTTCCTGA
AGTTACTCCAGGGTCCTGTCTTCCCATTGTTAGGTTTCCTTTCTGCTGTCCTGGCTTCACCTTTCGTAGCCAAGAAAGC
ACTAAATTTGGCTCTGAAATGTACATTTCAGTACAATCTCCCTGTCTTGGCAGCAGTGGTGAGGCTTTAGCAAATCGTT
TGTGATCCCAAATGAGCTTCAGCAGTTTGGAGTGAGGCCATGGCCCACATGTTTGATGCCACCAGCCTGATTTTTGTAA
TGAAAACTATATCAATAAGGTAAGTAGTCCCTTCTTAGGCATGCATTCTTTTCCAATTTTATGTAGTGGTCAGTCATTC
TTACAGTGATACAAATAAATATTTATGCTTTCATGTCTTTTAAGCTTTTAGCAACGTTAAAAGAAAAGGAGTCTGTAAT
GAATGAAATTGTCCCATTGAACTTATTTTTTCAGCTTTTGAGGATATATCGAGTGTACTTCTTAAAACTGAACATGTGC
ATTTGTTACATTTATATATGTTTCCACATATCACTTTGTAAAAGAATTAGAGAGAGCTTATGAAATAGAATTTAAGTTG
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TGTTCCTTAGGGAAAAACATGTTGTCTTGGGTACAAAATTCCAGCAATGTTTTTTTACCACATGAGACTTTATGGAAAAT
TCCAGATAACAGACCTTGCCCTAATGGTATGTGGGCAACAAATCCAGAAGTATCTATCATAGCATTCTTTCTTATAACA
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AGCATGGACAGTTCTGGTTATTAACTCACTCTTACATAAAACTTTTAGAACCAAAGGATTGGAGGGATGCTGACATTCC
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AAAACTCACTTTGAGGACCTGAAATGTTATATTCCACTTTGCTGATTGAGGTCAAGGACTATTGTTTGAACCTCAAACA
AGCAGGTGGTGAGGTTCACTTTCAACTGGTAATTACGAGAAAACGAACCATTATGCTAGATGCAAGGCCACCTCATCTT
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GGAGTTTTTATGCATACTGTTTCCTCTACACTGAATAATACCTCCAAACCCTACCGTCACTGCCTATTAACCCCCTACT
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Fig. 6.[61]

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CCCTTAAGAACACGATAAGAGTTTACCTTCTTCCTCCCTAGAAAATATACTGCTTTTTCTTTTATAATTAAAACAATAA
TATGAAAGTTAATCTTAGTCCAGAAGTCCTAATAGCCCAAGAATAGAAATGTCAAATTCATCTCATATCATTAGTTTGT
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CCAATTGAGCTTCTTATATTACAGATAAGGAAATTGAGGCACAGAGATATTAAGAAACTTTTCCGCAGGAAAAAAAATC
TGCTAGTTATTGAGCTGGGATTCAAATTCAGGTAGCTGATTCTAGTGTGCATGTGCTAACCACAGCACATACTGTGGCT
AACCATGCCTACACTCATGCTCAAAACTACGCTGTGATAGGTCTTATTAGCATCCCCATTTTACAGACGTGGAGATTGA
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Fig. 6.162

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GGTGTAGAGTGGTTAAGTGACCTGGCAAAGGTGGCAAAACTAGCATCACAGTCAGCCAGTCTGGTTCCAGAGGTAGGAA
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GGATTTAAACCTATTTTCATCTTGCAAGTATATATTTTAAAAGTGATTCATTTAAATATTCTGCTCAGCAATTAGCACA
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CCTTCATCTCTCCTTCCTTCCTTCCATTCTCCCTACTTCCTGACTGCCTATGTGCTTTCCTCCCTCCCCCCTTCCCTCC
CCCCTTCCTTCTTTCCTTCCTTCCTTCCTTCCCACCTTCCTTCCTTCTGCTCTCCCTCCTTTCCTCCATTTCTCCCTCT
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Fig. 6. 163

CTCCCTTATATTTCTTCAATTCTTATCATTTGAGTATCTAGTTAATGAGTTCTCATTTGATGTTATAGATTTTTTTAGC
TCTTCCTCACCTGTATTCCTTGGAGTAAATTTTATGAATCTTAGAAACAGAGTCTGGCTGATAAATTTATTGGTGCATC
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AATTAAAAAGGAAGTCACAAATTGAAAAAAAATTATATGCATAGACAAAGAGTTGATAAATTTAACATAGAAACAGGTTT
TACAAGACAATAAAAGGATAAATGTAATGATTTTCAATAGATATAAGACATTAATAGGTGGGCATGTAAAAAATGTCAA
GATAACATGAAAAAATTAGTTTTACTACTCATAAAAGACATGCAAGTGAAAACAACCAAGATGCCATCTCCAGTCTATC
AGATTTCTCTAAAAATGAGAAAGCGCTCATAACCAGTGAAGGAGAAAAGGATCTGCTTTCATCTTGGTTAAGATATACA
TTGCAACAATATTTATGGGAAATTCTTTGGCTATATCTATCACTATGAATTTATGCATACATTTTTCACCCAATAATTA
TACTTTTGAGAAGTTATCCTAAGGAATTAATCTAAAACTGAAATGATACATGTTCAAAATTTTCATTAAAGAAAACCAC
CAAACTGTACCATAATGGGAACTATTCATAAATATGTAAATTATGTCCATATGATAGAATACTAGGCACCCATTAAATC
ATGTTGTAAACAATATTTTTATTGAATTTTTTAATTGAAACCAAAAAAGAGCAAGATAAACTCAGTATACATAGCATGT
CATTTTTGCAAATGTGTATGCAAATGTGTAACATGTATATTTTTAAGATTGGAAGCTGGTACACTAAAAGTGAATGTTA
ATTCTGGGTTATGTTTTTCCCAAATTTTTGCCTGTTTCTGCCCTGCCTCTTTGATTTTAACATTGAGCATACATTTATA
GTTAAAGACTGTTAAAAGTGGCACTTAGATATAGCTTAAAAATAATTTTAATAGCACAGGGAAATTATTTTATTAATG
CTAGTTAAATTAGACGAAAATATAAAATTTTAATGGCAGAAATATTTCAATTTATAACATATGCATAGGAAACTGAAAA
GAGTAGTTAGCTCTAAATATGGGGATTATGGATAAATTTTATATTCTTCACTTTTTTATTTCAAAATTTTTATAATTAG
CTTTAGGATTTTTATAACCAGAACAAATTACATTTAAGACTCCCTTCTAAACTTATTTGCTTATTTTCAATCTCAAATT
GTGAATTTGTATGGCCTCTATATTTCCAATCTAATTTTACAGACATTAAATTTCTCTTGAAATTCAGTGAAAAATTAGC
AGAGTCGAAGTTACACTTCTGTATGGCATTTAAATTCCTCCTCCCAGAATACAGCCACTGTTCTACAGTACAGAGGAGT
CCTTCATATCACGATTTTCATTGTGTCTAAGAGGCATGTGTTTCCACTTTGTCATTATTTGGTCTAAAAGGATTTTTCT

Fig. 6.164

TCTATTTTGCTGCTGTTTTTGTACAAGACATATAAAGAAAGGTGTTAAGGGAGTGTTAAATATAATCAAGGTGTTAAGG
AGAAGAGCCCAGAGAGAGAACTCTTAATCTTTTCATCTGGTCCAGTTGTCAAAGTCTTTTGTCTTTAGCCATCACTTTT
TTAATCAGAGACATATAATCTATTAACTATGGGAACAAGAACCACAGATTTAGCCTGTGCTTATCTGGAAACCTCCCCT
AAAATACTACTGAACGACTCTTTTTTTTTTTATTCCTGTCTCTTTTTCTGTTGCAAACTGATTGTCCACTATAACCTGG
AGAGGGGACTCAATTTTTAGAAACTGTCTTATACACCTCTTAATTAGTTACCTATTGCTGCATATACCCAAAATTTAGC
AGCTTAAAACAATAATGAAATTTTATTATGCCCACAGTTTCTTTAGGTCTGTAGTAATTTAAGAGAAACTTAGCTGGGT
GATTATAGCTGACTCTCTCATGATATTACAGTCATATACATCAGGACTGCATTCTTCTGAATACTTGACTAGAGCTGGA
AGATTCACTTCCAAGATGGCTCCCTCACATGGCTGGCACATTAGTGCCTGCTGTTGGGGGGAGACTTCAATTCTTTGTT
GGGTGGTCTTCTCCATGGGGCAGCTTAAGGATCTTCATGGTTTGTGGGCTGTTTCACAAAGTGTGCGTGATCCATAGGA
GATTAAGGTGGAAGCTGCAATGTCTTTTATGATAGAGCCTCAAAAGTCACACAATGTCATTCCCAACGTATCCTTTTTT
TCACATAGGTCAGCCCTACGCAATGTGAGAGTGAGGAAGGGACTACAGAAAAGAATGAATTCCCAGAGATGAGTAGGAT
CATATTGGGTGCCATCTTGGAGGCTGGCTACCAGTGAAGAAGACTGCTTTCTGGCCCTCAGTGATTTACTTCCATCTCA
CATGCAAAACATACTCCCTTTATTGTGGTTCCCAAGAGTCTCATTCTATTATGGCATCAGCTCGAAGTCCAAGATCTTA
TCATCTAAGCCAGGTGTACTAGTTTTCTTTGCTGTCTAACAAATAATTTAAAAAATGTAGCAGCTTAAGATAACACCCA
TTTACATGTCACACCTCTGTAGGTCAGAAGTGTGGGCAAAGGGTAGCTGGGTTCTTAGCTCGAGGTTGAAACCAAGGTG
TTAGCCATGGCTGCGGTCCTTATGTGGAGCTGAGGGTTCTCTTTCAAGCTCATCCAGGTTGTTGGCAGAATTCAGTCAC
TTGTGACTGACAGACTGGTGTTCCTAGAGACCACCTGCCATTCTGTACTGCGTAACCCTCTTCACAATATGGCAGTTTA
CTCCTTCAAGGCCAACAGAAGAATCTTCCTACTGCTTGGGTTCTCTGACTTCATTTGACCCTGACCTCTAAACCCAGAT
TTAAAGGTTTTATATGATTATATCTGGCCCACCAGCAAAATAGTCCTTTTTATTAACTCAAAGCCAGCTGTTTAGTAAC
CTTAATTATATCTGAAAAAAACAGAATCAAAGGAGTGGTATCTCGTAGTATTCACAGATTCTATCCACACTTCAGAAGT
GGAAATTACACAAGGCCTGAACATGAGGAGATGGGAATCTTAGAATCTGTCTAACATACCAGGTCTATGTGCAAATAAG
ATCAATCTCTGTTAGTTTACTCTTATGGGGTAATTAATTACCCCAAAACTTAGTGGCTTAAAACAATAGCATAACTAAC
TTCTCAGTTTCTATGGGTCAAGAATTCAGAAGCATCTTTCTTCAGTGGTTATGGCTTTGGCACTCATATGAAGTTGCAG
TTAAGAAGTCAGTCAGGGTGATAGAAATATAAAGGCTTGACTGGGGCTGGAGGATGTGCTTCTAAGGTGATTTACTCAC
ATGATCAAGTTGGTATTGGCTGTTGCAGGCAGGTCTCATTTCTTCCGCAAATGAAATGCTCTCCAGGCTGCATGAGTGT
CTTCATAACATGGTTGCTGATTTACATTGGATGGAGAGATCAAAGAGGATGAGAGGAAAGCAGCAATGTGTTATAGGAC
CCAACTTCAAAAGTTACACATCATTGCTTCTGCCCTATTCTGTTGGGCAATCCTGATGCAATATAGGAGGCACCATGAA
TAGCAGGAGGCAAATATCATCTGGGCTTTCTTGTTAACCTCAGTTAACAACAACAAAACAAATATTTAAAAAACTAAAG
GAAAGAGATTTCTGGAATATTTCCATTCTCTGTCATGCCCACTGCCATAGTCTACTCAAGTTCTCATCTCCTATTGGAC
CTCTGCGGTAGCTTTATAAGTCCTGTCCTTGTTTCCAATGTCATCCCTTTCAAAACGCTTCAAATAGCCTCCTTAAAGC
CATGACTCCTCCCACGGTTATAACCCTGCAGGTAAAGGCACTTGAGGACATAATCTGTACCATCTCTGGCGTCATCCCT
TTCATTCCCCTTCACCTCCCCCAACTATTTTTGTATTTCAGTCATGCTGGACCTCTTAATGTTTCCCCAAAACATCCTA
AGCAATTTCACGTCTCTATGCCTTTTCTTTTCCTGTTCTCTTTGCCTGAAATAAGAACGACAGATAATAGTGACTATGC
AAGTCCTAAAATCGAGGCAGCCACAGTGCTAAATCACACATTGGATAGTTCAATTTTATTCTTATAACAACCCTTTGTG
GTAAATCTTATTACCATGCTCATTTTACAAATGGGCACTCCTAACCCGATGTGCATAATGATTAAAAACATAGACTCAA
GAAAAGTTTGGAGGTTCTGGCGTAACCCTCTTTGTTATTCTCTATGCAACTAGCTTAAAAATAACAACTTTGATTATTC
TAATCTATTAACCAGTTTTCAGTGTGTCTGACCAAATCAGTTGAATTTTATTTTTTTAAATATGTGTGTGTGTGCATGT
GTCAGTATGTACGTGTTAAAGAGTAGAGGGAGGAAGTTGAAAATCTGACATTTGATTTATTTCTCCTGGGACAAAGGAA
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CCCCTCTTCCTTTCCAAACTTCTTTCCAGTCTCATGACCAGCTTATTGGCTTCATATTACCAAGTAAATATATCAACTG
CCCATATGTATAGTACCCAGACAATGAGAACACTGCCTAAATATTGTGTTTTCAATACTACTTGCTTTTAGCAACAAAT
CATAGCACCCAGCCTAACTTTAAAAAATTCCATGTCATCCAACTTCAAACTATAACAAAAATTGAGTTTTACAGAAGAGT
TTCAAAACATTTTTATGCATTAGACACTTCAGAAATTATCTCTTAAAAACAAAACAAAATAAATAAGGGACAAAAACAA
TAGAAAACGTCAGGCACCAAACTGTGAACTCTCTTTTGTCCATGTATGATTTAAAAAAAAAAAAACTCACTGATGGTAAT
TTAAGAAATTTGACCTTTAGGAAAATCAGGCTGAGGCTCATCTTTGTACTTTTTAAAACAAGTCAATAAAACATATAAA
TAATAGAATGAGTGCCGCTCACAGGGGACACTCATATTTTCTAGGAGCCTGTGGATATAAACAATGAATTTCATGAAC
ACAGAGCACATTTCTACATGATCTCCTTTTCTTTACAAGTTATTTATGTTTTACTGGAAATATTAAAACTTCATGTCTC
TTTGTTGCTCTTGTTTCTTTAAATTTTTATTTTTATCATCAGTCATTGTTCAATAACTCAAAATCATATAAAATAAAAC
AGTTTCAACATCATTCCGTACTCAAGATGATCTCTATTAAATAATGATTTTTTTCTTGAAAAGCATTCTATAATTTTT
TTTAGTTTTCCAGATGACTATTTAAACCAAATGTTTAAATGAATGATTAATGAGTCTTTTATCTTATTTATCCCACCTG
CCTACTGCAAAAAGACATGAATTTAATACATTTTAGATTTTAAACAGTTGTATATCTGTAAAAAAACAATTTTTACAGG
TTTCATATTCTGGACACTAGAAAACACTTCCTCCTCACAGCGCCGAGTTGTATGAAAAGGCAGCTCAATTGTCTTTCCT
TGAGCAGCCTTGTGTGCCTGGGTATTCCCTTCCATCTCTCACCCATGGAGTTCATATTCCTGTTGTCATGTCACCACCT
CTGTCAACGAAATAGGTTCTGCCATTTCTCAAATTTTATGGAGAATTATCCTGATGAGTTAAGGCAGAGGTTGAAAACT
TGCAGCCCTTACCCCGGTTACGGATTTTATTTTTAAAACCCAGTGTTTTTGAACACTTTACAGCTGATATTTCTATTCT
AATGGATGCATTATCTTCTTTTAAAAATACTTCCTATATTAGCCCTCTTAAGGGGTATGAAGTGGTGTCTCATTGTAGT
TTTGATGTGCTTTACTCACTCTTAATACCGTTATTATAGGTTATTTATTGTTAAAACTGTGATATATTTTCCTTTATTT
TCTACATATACATAGATAACATGCAGTTTGATAATCTGCATTTAATCGCATTAGAAAGGTATTTTGCTATGTCATTAAA
ATTTTTTTGTGAACATCATTTTTAGTGGCTGCAAAAACTTACATAAAGACAGACATGTCCTAATATATGTGTGTGTATA
TTTTAACTAAATATTGGCACACCAAAAATAATGGATTTAATGCAACTAGCAATGATAATGCTCAAACAACAGGAGTGCT

Fig. 6.165

ATGAAGACAATGAAACAGGTGAGTTGTCAGAACACCTTCTGTCATTACATTAACTTTTTTAAGCACTAAATTCACTTGA
AATTACAATAAACAAGATAAGTACAAGCATTAGATAATTTGACCTATCAGATATCCCAAAGGCATAAACTATCACTAGA
GGGTCTTTTCTTTATGCAATTAGGTGACCAGTGACTCTTATTTTTTAAAGCTTTTAAAATAAAAAAATAGCTTTAGGGA
TAAAACCAGCCAGATTTTTATCGTCAAAAACATGTTCACAGTTCAGCTAAGCCTTGCTTTCTTTTGAAGAATTGCTCTT
TATAATGCTCACCAATGGATTTAGAAATATCCCATTGTAATCATGAAGATCTCAATTATATGTTATGAAATAAGAAACA
TAATGAAATACTTGAACCTTCCTGCCCCGCATAGCAGATATATTTTTGTATTTATGATATTCACAGTAAAAAAGCTGTG
CTGAGAAGCTCAAAATAGTATTTGATCTTAGTGGTTTTGGAGCAGAGGAATTGGCATTGCCATGATTTTTCTATAACTT
GGCGCAGGCCAATGGTTTCAGGCAGGACTTTCAGAGCAGACTTCTGTCAAATTTGCCATATGGAGTTACTGGGCTCAGA
TTATAGAGGTTTTCCTGGTTGTTTGCCTAATTTGGGAATTATCAGTGCTCTCCAGGATTTTCTCCAAAAACAAAGAACA
TGTTCAGAAGAAACTGACACTTCAGGAAGAAAAGTAGTTTGATATTTGAGAGGTTTAGATGGTTTCTAATATTTCTAAA
AGGTACCATGACTCGTGAAGAATTTGACTGGTAAAAAAGAAAGGTTCCTGTTATCCCTGGAACACAGCTAACAGTCTGG
CATGTAGCCTTACCTGAACTGGTACCCTCCCCGTACACCTCACACTTAGCATTCTGCACAGCCAAAGACCCTGTCAGTT
ATTGGCCAGCCTGAAGAATGAGGTTGAATTCAACTCAGGGACTGAAATCTACTTTATAGCATTTTAAAATATTGTAGAT
CCAACCTTGAGTTGACCTACTCCGCAGGTTCATGGGTGATTAGGTGTGAGATACAATGGGAATGAATATCTGAAAACCC
TTTGCAGAAGAGAAGGAGGAGGAGGAAGAATAAATCCAATTTTTACAGGTTTCATATTCTGGACACTAGAAAACACTTC
CTCCTCACAGCGCCGAGTTGTATGAAAAGGCAGCTCAATTGTCTTTCCTTGAGCAGCCTTGTGTGCCTGGGTATTCCCT
TCCATCTCTCACCCATGGAGTTCATATTCCTGTTCTCATGTCACCACCTCTGTCAACGAAATAGGCTCTGCCATTTCTC
AAATTTTATGGAGAATTATCCTGATGAGTTAAGGCAGAGGTTGAAAACTCGCAGCCCTTGCTCCAATTACAGATTTTAT
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AATTCACTGGGGTCTCCATGTGACGTTTCAGCCTTTGCACCCTACTTGTCCGCTGCAGGCCCAGGAGCTCTCTACTCCT
CATATGCATTACAGGACCCCATGCACATTGAAGATGTTTTCTCAGGCAGAGCAGACACCAGGCAAATATCTTCTGGTCT
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TGTAATTCTAACTCATCAGTCCAGCAGTGCTGAAAATGCAGAATTTTGTCCTTATCAAAAGCCAATTAATGCTAACCTT
TTCAGTGTGTGTCCAGTACCACCATCGGTAAGGTAAATAAACAGGCAGTTTTGCATATAATCTAGTGATTTTTAATTTA
GTTTTTTCCCTTACTCATCCTAAAAAAAAGCAGGGAGTCATTGGGCTTTTGTATCTATTTTAGCTGAGTTATTATTTTAA
CCCCTATCTTTAGTGCAGAATTAAAATATCACCCTTCTATTTGCTTTTATTCCTAATATCTGAATGGTTTCAGCTTGAC
GCAATTAGCATGTAGAGCATGCTCACCTACTGTAATCTCCCAGATGAGTCACTCAGCTCCAGCACATAACTGCTTCTGG
GATCCTGTTGGCTAACCAGGCTTTGAAACCCAGGGAAAAGAACTGTGTGTTTTCTTCAGAAGTAATCCATAAATGAGAA
ATTACCATAAATAGGCAGTTTTATTCACTCTAAGTGATTTTTACATAAAAACTGACAAGGCCGTAAATTGTATATAGAA
TCAGGTTCCATATCCACACAGCTAATGCACACATGAATTATTCGATGAAATTTGCTTCAGTAGTATTATCTGAAATTGG
TTTCAGACAATACATTAAATGTTACACATTATGGTAAATTTACTGTATTTTTGTTCCAGCCACGTCTTGTGTTGGCAAA
TCTGCTAAATTTAAGTTTGCATTGTATTCACATGTGTTGCTATTTCCTGAGATGTCCCAATATATTTTAGCCATCAAAC
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ACTGCCATTCTATCTGTGATTGCTGTTCATGGAAGAAGTAAAAATGAGCTGGGACAATATATAGGTGGCGCCAGGGATG
GCTCCTTGAAGGAAGTGATCTCACAGCTCAGACCTGAAAGCTGAGTGGAAGTTACCAGAGGAAAAGAGAAAGGGAATGC
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GGGTCAGGAGTCAGGTGCGTATCAGAGTCTCAAGAGGCTGACATCAGGTGTTTGCAGATGGCATTCATGTCTGGGGCTC
AGGGTCCTCTTTCAAGCTCCTTCAAGTTGTTGGCAGAATTTAGTTGCTGTGGTTTTAGGACCGAGGTTCCCACTTTCTT
GCTGATCACCAGCCTGGGGCAGCTCTCAGCCCCTTGAGGCCACCTGCAGTTCCTCATCCTTTGGCAGATCCTCTCACAA
CGTGGCCACTTACTTCAAGACCAGCAAGAGAATGTTCCTCTTCAGGAAGGGCCCAACTCCTCCTTATGGATGTTCTTCT
GATTCAGTCAGGCCCACCCAAAATAATTGCCATTTTTGGTTAACTCAAAAATCAACTGATTTGAGATCCTTAATTACAT
CAGCAAAATCCCTCCACCTTCTCCATACATAATAACCTAATCATGGGAGTGACTGTTCATCCTGTTCGTGATTAGGCTC
ACATATAAGAGAAGGAGATCATATAGGACATGCATGTAGAGGATAGAATCTTGGGGGCCATCTTAAAATTGTGCCTACC
ACAAGGAAGGATTTTCCAGGCAGGAAGAACCAGATATATGAAGGAGGCCTAAATGACAGGATAGCAATTTCCAGTGTCT
GAGGCTAAGGCTGGAATGCTGACCTGGCCAGATCATGCACTGCTAGGTTAATGGCCTATGCATACTCTATTATGCAAT
TATTAAAATACCTGTTTACAAAGAATATTTGAAATATTAAAAAATGGAAAACTGCATACCGTAAATATTAAATGGGAA
AATCAGGATATGTGTTTATATAACATAGTATCAGTTTTGGAAATATATATGCATATCTTAAAAGACTAGAAGTAAATAC
ATCCATCTGTTAGCCATGGATATCTCTGGGTGAGAAAGTATGGATGAGTTAGAGATTCTTTGTCCCTTTTCTTAGTTCT
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TGAATTTGAGCAGTCTCTTTGAAGGAATCAAATTTTGTACTCCAGGTGTTTGAGAGGTTGAAAAACAAGCTCAGGAATT
AGCTGCGCATGATCTGTACAAACAAGGGATCAGTCAGTGGAAATGAGTCCAGCTTCAGCCAAGGGTGGTGAGTATAA
CAGACCTACTCTGATATGTTTCATCACTAAACATTGAGCAAGAAATCTAGTGTTTAAATTATGGAGGCATTAAAGCAAT
TAACCTCCCTCTACCTCAATTTTCTTATATCTAAAAATTGAGGGGAGCAAAATTCCATTCTTGTCACAGGTCTCAATTG
CAAGTAGCATGCAAGCAAGTAGGCTGAACTCTTGTCTGTCAATCTAGCTCCTTCCTGGATGGGGTATTTAATCAGCTCT
CCTTCCCAAGCACCCAGTGGACTTGAAGCTCACAGTTAGACAGAGATGGCTAGAGGTGCCAAGTGCTGTTTATCATAGG
TGACATCACCTGTGTGCTCAGAGTTATAAACATAGGCAGCTTGCCTACTGCCAAATCCATTACAACTTTCAGAGGTGTT

Fig. 6.166

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CACTTAGGAAGAGGAGGAAAGAGTTTTTTGAGAACCAGATGGGAACTTAGAAATAAGTAACATTGACAACCAAATCCTT
CAGACACATCTTCAAGTTTCAAGAGAGCAGTTAACAAAAAAGAAAAATGTATTTTTAGAAATTGTTTGGGAGGTGAAAG
GATTTTTCTGGGGGGGAGTAGGGTAGGGAATGGATAGAGATGATTTTTTTTTGGCTCTGCAATATCTCAGAGATCTTTG
AGTTGAAATATAACAAAATGTATTTCCTCTCTCTTGCTCCAAAAACTGAAAAGCAAGTGCACTAGAAAATGTCAGAGTT
GGCCACTGAACAGACCATATTTCTAATTCATGGCAAGATGGCTAAAATAAAAATTCTTGAAATACGTATTTTAAAATTA
GATTTTGACCTTTGATAGGGCTTATCATGCTATTTATATTGTAGCAGAAGGAACTGATTTTTTAGTAGTTGCCTCATCA
AAGACATGACTCACATTTCCTTAAGCATTTACACCATATGTTTCCTGAGGTAAAAAAATAAATAAATAAATTTTATATT
TTTTATACAATTAGTGAAATGTGAGCAAAAAATATAAGAAATCATTTAAAAAATATTTTGGGGAGGATGTGGTCTAGAA
ATGTTGTTTTCTAGAAATTGAAATTATGTTAGCTATTTGGCCAGTGTAGCCATTAGGTACATTTATGAATATAGATGAG
AAGTAATCATTGTGATGCTAGTTTTATCATTAAGGAAAAAAGCAGAATTCAAAAGTTTAGCAATGAAACACTGGTACTA
CTGAAACTCACTTTATATTTGAAAATGTGAATGTGAGTTTAAAAAAGCTTTTGTTAGACTAACTGAAAAGAATCATTTT
AACATTTAAAAAGAAAAATATTATTTTAAGTGACATTTGGTGACATTATTATATGTGTACAAGCTGTGACTATAAGATA
ATGAGACAAATTATTATGTGGCCAGTGGAAATTAAATTTATCTTTATGATTACTAGCAACAGGGACTATTTTATTCAAC
TATCTTTACATAATACAAAATTATTGTAAAATTTCTAGGAAATTCATATTCCACTTAAAATATGAGGAAGTACATTTGG
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TTTTATTTCATTGGTACTTTTTCTAGGGCAGTAGATTTAGGAATAGAAAACAGACTCTACACATGGAATACACTAAAGT
CTTGAGGAGGTGTGATATTTCCCTAAGTTTGGATAGGTGGTATCATCCTTTCAGTCACTCTTGACTCTGCATTAGACTC
TCTACCTAATGATCCCCAAGTCATTTCACTTCTGCCTTCTTAACATTTTGAAGATGTCCTCTAAAATTTCAAGTCCCAC
CATCTCTGCCTCCGAACTCACCTCCCTGACTTTCAGTCTTGAAGCACCTTAACAACCCCTCCCCTCCTGTGCATGTTCC
ACCTGGAAACCAGGGTGATCACTCTGAAATGCCAATCTGATCAGTCATTTCCCTACTTTTGAACCTTCAATAACTTCCT
AGCACCTATTCCTAGAAACCATTCCAACATATCACCTAGGCAAAGGAAACCCTTTTCTTACCTGGCTCCTACTTATCCC
TCCAGATTCACCTTTTGACACCTCCCACTTCATCTCTCTGCAGCCCCCTTCCTGCACACACGGAGTTTCAGTCTTGC
CAAACTCTTCGCAGCTTCTTGAGTGGACCTTGCTCTCTGCACTACAATCTCTTTCTGCCTAGAGAGTCCTTTACCTCCC
CCAATCTGGTTAACTTCTACTGTTTATTTAAGTCTTAACTCTAAGTAGATGTCATTTTTGAGGATGCTCTCTCTGTTCC
CACTTTGGAACTTGGGCTTCTCTGTTGTCCTGGGTGTCCCGTGTATTCCTCTGTCATAGCCCTCATCTCATTGAATTGT
AATCATGTCTTCACTAGTGTTTCCTTTCTAGCATGATAGATTTTTGTATTGCCAGTGACCCCAACAAATTCCCCAGGTC
CTAGCCCTGTGACTGGAACACAGTAGGCATTTAATAATAGTTGACAAATGAGTTAGTATATGTAATGAAAAATCTACCT
GCCTGAGCATTAGGAAATATAAAATTAATCTAAACTTTGCCACTTTCTGTAATATCTGTGTGTCAGTCAGGAATGCTTT
TGGCTGCAAGAAAACCCCACTGACAGTCTTTTAAAACAAATAAGGAAGTCTAAAAGGAGTCAGATGCTGGCCTTGGTTC
AATACAATGTTAGAGGCAGCATCTCAGTGCTTCTCTAATGGTCACAAAATGACTACAGTAACTCCTTCCCTCACATCCA
CTTCCAGTTCAAGAAGCAGAATGAGAGACAGCAATAGCCACCCTGTTTCTTCATCAAGAAAGCAAAGATATCTTAGAAG
TCCCCACTGTCCTTCATTTACTTCTCATGGGCTAGAATTATCTCAGTGGCCACCCCAAGTGCAAGAGAAACTAGGGAAA
TTAGTTTTTAGCATTCCCACACTCTATGTTGAAAATGAGTAAGAGACAATAATTTGGGAATAGGTGTGGGATTAGTCA
GCCAACAGTGTTTGCTACACCTTGSGACAAAACATTTTACCTCTCTTACTCTTTGATTCTTTCATTGAGTGAAGAATTA
GAATAAGTAATCTTTCTTCAAGAATCAATTTTCTCTCATCTATAACTCCCATATATTCTTATTTCCAAAATGGAATTAC
CTTCTACACCAGACAGTTACTTCAAGCTTTTCCCCTGAATATTTCACATTTCCAGAAAGCACTATATTACAAAGAAGCC
CTTCCTTTTCAAGAACTCTATTTGCCATAATGTTCTCCCTAACAAGGGCTGAACAGTACGTGCCCAGATTCCCTGCCTC
TGACCTCACCACACCCTCTTCTGTGTGGCGGTCTCTCAAATTGTTTAGGACAATTATCATGACTTCTCTAAATCTCCCT
TTTTCCAGGCTCAAATTCCCTATGCTTCAGTAGTTTTTCAGATAAGCCCCTTCTGTCCCTTCACAACCCTGGTCAGCCA
ACACCAGACATACTCTCTTCATTTACTTATGATTCATTGAACAGATAATCACTGAGTTCCTGCCATGTGTCAGCACTGC
TCTAGGAGCTGGTGATATGGTGAGAAACAAGGAAGGGAAGGCCCCTCCTCTCATGAGGCTTACAGAGGAGCAAAGGAAT
CAGATGATAAGATAAGCGTTACACACACACACACACAGAGGATCATTTCAGAGAGTGATTCGTGCCAAGATGAAAAT
GACAAAGGGAGATAGGAGAGTGACAAGGGATGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGGCAGTGCACGTGTGA
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ATGCTCAGAGGGGGAAAGTGTTGGAAGATGGCCTGATAATGCACATGGGGAGCCCCCCACCACAAGCTCATGTAGGACA
GGACAAGAACTCAGATTGTATTCTAAATGACAGTATTTCAATACCCATCTTAAAATGTGTCTGGACAAAACTTAACACT
TAGCTACACATATAGCCTGACTGGAGCAGTCCAGTTTATAACCTCTCCTTGTAGTGGGCACTGTATTTCTATGGATGCA
AGCTACAGTTATATGACTTTTGTGGAAGCCATGCTATCCCCACAGTTCCACTGTGATTTCCTATATGCTCTTTGATGCA
CACTGCTCACATGCATGAATATTACTTGCACAAAGAGAGAGAGTAGGTGGATGTGATGCCAGGAGCTATTAGAATGATT
CACATTACACTTCCACCATATGAAAAATAATGTACAGTCTAAAGCAACACTGCATCATATCATCGAGCAGCTTGGGAAA
GCCTAAATTACATATTTGAAGTTCAGTCAGTCATTGAAATTAGATAATCTGGATGGTCCTGAAAGTGTCTTTCTTGAGA
GCTTTCCTCCAAAAGCAAAATTGAATGGTAAGGAAAAAAATGAAGTACAACAAAAATGACAGCAAAAAACCTGAAAAAC
AAAAAACAAAAAACAAACAAACAAAAAACCCCAAAAACTGAGTAATATCAACCAGGCAGGAGTTGAAGGAGAGAAGCCC
AACTGTTGGTAGATAACTTGGGATGAGCCAATTAAATATTGGTCTTCTCTCTCTCAGAGGCCCTAATTGCATTCCCTCC
AGCAGCCACAGGCCTGGGGTTTTCCCCAGGCCTGTGGCATTCAGACATCAGAGCTGAGAACTCCAGTGACCACCAGGCT
GCATTTTAATGCTAAGCTGGGAGCATTGATACTCTCCAAGACAGCCTTGCAAGCTCCATCTTCCACTGACTCAACGTTC
AGTGGAACGTTCAGTGTGTTTCATCTGCTGTTCAGTCATGCAGGCAGCAGGGATCCTGAGGGCTCTCCTCCCCAATATG
GTACCAAGTTTTGATTCATCCTTTTCCTCTTCCCTATCAGCAGGTATTCTAGAAGACAGTTTTATGATGACATTCACAG
CTAGCATTAAAACCAGATAAAAATAAAACAGGATGTCAATAAAGAACTAATGATGTGTGTCTGTGTAAATGTATAATAA
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Fig. 6.167

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TAATAAAAAGAATAGACATCTGTTTGTCAAGGTACCATTTAAGGCCTTGGGGCTTTAAATGAAACTGCTTTATAAATGC
ACCAACATATGTGACTGTTTGCCTACGTCTTTCATCCATGAAAATTGCCCTGTAGACAGTTGCCCTTTGAGGTGAAAAGC
ATTCTATAGGCACTAAATATTATGCTAGAACACCTCTAAGTGCTAATCTAACTACCTTCATGACGGAAGTTTCAGGAGA
ACAATTTAAGGGTCATGGAAAAAAAGTGTTAAACGAGTTGGTAATCTGACATTTCCTAAAAACTAGATTCAAAACCGAA
GGGAAAGTAGAGAGAGGAATCAAAACTAGCCTAGTAGGATGGAAGTAGGCCTCCCTCCTTGGAGGTGTGAAAATGCCTG
GAATAATTTAGGCAAGAATGCCAACTGTCAGTGATGACCTTGGTTTAGGGGCAGTTATAGCATGGTGGAAACACCCCTG
GACTGGAGATCCGGCATCCCTTATGGGGATATGAAAGAAATTTGGCAGGTCTCTTAACCTCTTTGACATTCCTTTCTTC
TTTGTTAGGTAAGAAGCTGGTGCTGGTTCAGTGTAATTTGAACTGTGTTCTGCATGACTGTAGGGTTCTGCTTAGTGCC
ACGTGGAGGTGGGGATGCCTTGGGTGAAGATGAAGGAGGTGTAGTAAGCAGGTGGGGCTCTGCACTCTTTTTACTCTCC
TTCTTCCAGTATCATCAGAACAACTCCACTTTTTTGTCTTTATACACTGAGTTATTTATAAAATCCTAGTTTTTTTAAG
TGTCAAACAATGCTTGATTATGAAATGATAGCATTATTCTAGCATTCTAACTAAAGTGTGTAAATTATAAACAACTATC
TTTAATGGTCAACAAATTGTCATCATTACTGAGCAGTGTTGTTAGTAACACCAAGTAATACAATTAATGAAATCCTTGA
AAGTAGAACCTATCAGTGAGCCATGCATACAACTGATAAGCTGTGGATAATGTATAGAAGAATGAGTTCATTTGGTAAT
TATTTGTACTTTTCAAAATATTGGTGAACTTTCATCTTTAAGGAGCAATGGTGTCCTAGTTATTCTTGCCACAACTGCA
TGGAAACTCTCGAATCTGTTAGTATGTCCTTGGGCACAAAAACAGTATTTTTTGGAACTGCTCAAAAATTATGTAGTTT
TGTAGACATTTAGGGAAAAATCTCATAATTTATGTTAAACACTCAGTTGTCTTGGTGTTGATTGAAAGCAAATAATACG
TGCTTAACATCCTGCTAGTTGCTGAGGGAGATTCATTCATCATTTTAGAAAATGTTTCTGGGCATCAGTCATATAATAA
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```

Fig. 6.168

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Fig. 6.1169

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Fig. 6.170

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GCCAATGTGCCTGGCCCTAATTTTCATTTATTATACTTCCCTTACTATAGATTCAATACAAGGGAATGAAAGCTTAGAC
GTTCTTGGCATATTTTATGAACTCCTATAAACCCCAACCAGGTTGTGAGTAGTTATCGCTTACTGAGACCTTAAACACA
AACAATATTTTAATGTTCATTTATTATCCTGAATAAGATACATAGATATATATTGGGAGTTGAGCGCTACCATGTCAAG
AGGAGCAAGCCCATAGAATGATTTTTCTTTCATTTTTTTGCAAAACAACTGCAATGTCCCATTAAGACTTTTCCTGCAT
GTTCGTGTCAGATGAAGGAAAGTATTTTCCACTGATACCCCAGGTTGTTAGCTAGTCCCACGACTGCTCTTCAAGAAAC
ACAAGAGACCTATCTGAATGTGGGTAAAAGGATTTCATGTTGCTTTTTACTTAGAAACCCAGATTGTTTGTAATAAGCT
CTGTCTGATTCCTTCAACACTTAGCATCCAGGCAGTTTTCTTCCTCTAAACTTCAAAAAATAGGAGAGAATTGTAAGGA
AAATGATAAAAGCATCTGAAGTTCAGCAAAAGTGAGCCTCATCTGAGCTGACTTGTCCTGATTTGGGGATAGCCTGAGG
AGAGTGTGTTCACGCACCTTGCAGAGTAGCCCTTAGCAGCCACTTGAATAAAAACGAATGAAAAGATAAACAACAGGAG
TGGGCAGAAGTTCGAAATGGAGGATGTGTGTGAAGAAATAATAACATTGTCTTGAAGATTTTAGAAAATGGGAGAGGCA
AGAAAAAGAAACATAGAGATGAATCCTGGAAAAGGTATTTAAATCTTGGTTGATTGTTCATAAAGCCCCCACCCATAGC
ATCTTCATCATCCCTGTGGGATGCAAATATTGTTGTTTTCATCTTACACACACGGACATTGTGTTTCTGACAGGCCAAG
GGGCTGTTCAAGGTCAAATGACTGTTGAAGGCAAAACCACTAGTATGCAGCAGTAGCAGAAAATGAATGCATGGCTTCT

Fig. 6. 171

GATGATAAATCCTATGGTCTTTTCAAAACGCCTCTGTTGTCTCTAGGGAAATTGCCTCAAATTAGAAATCAACACATTC
TACTGCAGTTCTAATTTGCACACGCTTGCCTGAAAACTTTCTAATTTCTGGTTCTTTTTTACCACCTCCTCTCCCACTT
AATTTCATAGCTACAGAGGGGAAAGATAGCTACTGTCCAAATAGTCTTGCTAAAGGACCTCATTTTCAAAATTCTTTTT
CCCCCTTCAAGTCTTCATTTGACTCACGGTGTAATCACATTAGTGAGGAAGATTTTGTACAGTCATACCTAGTGCCAAA
TATCAAGTTTTCCTCCAAAAAGCAAAATAAAATGATGGGCAGACCACCTGCTTTCCATAAAATTGTACCAAAATCAAGG
AGGACAACTCTACGTTATCACGTAGAAATAACATCAGCATTTATGAGTAATAGGTCTGTTTGATGGGCAATGCTGTGGT
GTGTAAAATTCTGATGACAACATGTTGAAAAAAATGTCTGGAAGAATGGGAGAAAAAAAGTGAAAGTCTATGGAAAGTA
GTAGATTTGGAGTCCTTTTGAAGAAAACATGCAAGAAGCTGCCAGATAACTGCTAGATAAGCAGCTATTGACCATTTAT
GAACTCAGTTTTACCAGGCGTTGAGAAGGCGAGAATACTCAAAATAAAACCTGCATAGATTCCACCCTTAGGAAACTTA
CAGTCTGGCAGATTCAAAACAATTTCACTTCACTTTAAGTAATTGTAAAATTGCCTCCATAAAAGCATAGCCTGGATGA
AAACATTTTTCAATATTATTTACTCATTTTAATGCACTCAAATAAACTTTATACCAAGTAACTTCTTTAAAGATCTACC
AATATAAGGCACGTATAGATCAAAGTAGACAATGTCAGCCCATATAATCCCCTTCTCCAGGTCCATTTTGATTGCTTGT
GTCAACATTGGAGACCACAGTAAATTACTAATGGTCCTTAAAATTCCCCCCCTATTTATCTCTTTCTTTTCAATTTTTC
TCCTTCACTCCTGCTACTCTATTTCTTGTGTTTTTTTTTATCTACTTTTTTTTTTGCTCTTCCCCAGGGAAACTAGA
GGAAAATTCTTCATCCTGATAAGAGTTTTTGGCACCAGAAAGGATATTTTACCTGTAGTCTGACCTACATGGAGTGAGG
GAGCCTGAACATGTTTCCATGAGTGTTCTAGTTCAATGGGCAGAAAGATCCATCAGGAGAAAGCCTTTGTCACTGATGG
CTTTGAAATTGTTCAAATGACTTTGCAGTTCACAACACGTGAGCTTAATCATTGGTTTGGCTCTTCTGAGGAGCTGAGG
GAGACAAAGAAGCAAAGTTATTTTCCCAGAGGTAGTGAGGCCCAAGATGCTCAGAAAAGATTTAATTGAAAGATGCCAT
TGCTGAAATTTCTTTTATAATGTTCTGCCTGTGGCTTTCCAATAGTAATAGTAGTAGCAGTTGTTGTAATAATAGAGAT
AGTAGTTGTATTGATAGCAATAATAATAGCTACTATTTATTGAGCACTGACTGTGTTTCAGGCATGGGTCCATGCACTT
TACATATATTAACTCATTTAAACCTCCCAACAACCTAACAAGGTAAAGATGATTATTATCCCATTTCACAACTGAGCAA
ACTGGGACATGAGATTTAAGGAACGTGCCCAAGCTCACACAGCTAGGCGATCACCAATTATACATTTCCTCTCTGCAAG
AGAACAGTCTATTTTTCAAAAACATGCTGATAAAACAGCAATCCCATTTGCTTCTAAGGACAAACCCTGCCATGCCTGA
GTGTTGCCATTCTTAAACTCCAGGCCAGGGCTTGCCTTTGGATGCATGTGAGCTTCACTGGGCTCTCTGCTCAATGTCC
CTCCTCAGAGTGTCTCCCTAACCACCTCTCTTGATCTCTCTCCACCCCTTTATTCCACTTGGGTTTTCTTCAGCACACT
TAGGGCTACCTGACAGTGTTAGCAAGGCTACAAAGCAAAACACCAGATGGGGTGGCTTAAATAACAATCATTTATTTTC
TCACAGTTCTGGAGGCTGGAAGCTTGAGATCAGTGTGCCAGCTTGGTTCAGTGCTGGTTATGGGTCTCTTCTTGGCCTG
CAGACACAGACACCTTTTCACTATGTCCTCACATGGCCTTTCTGTGTGTGTGGGGGGGGGGTGTGGGTGTGTGCGC
GCGCGCGCGTGCCTGCATGCCTGTGTAGGGAGAAAGAGAAAGCTCTTTGCCTCTTCTTATAAGGTCAGATTAGGACCCT
ACCCTTATGACCACATTTAAACTTATTTACTGACTAAAAGCCCTATCTTCAGATATAGTGACATTGAGAGTTCAACATA
TGAACATATGAGCTGATAGGAATATATGAATTGATGGGAACATATCAACACTTCAACATATGAATTGGGGTGAGGGACG
CAATACAGTCCATAGCGGTGACCGAGCATATATTTAATAGGTCATTAATGTGTTTACTATGTGTCTTTCCCACTCCTCC
CGGTATTAGTAATCTAAATACCAGTAAGGATTTGAGGTTCTTCTTTGTTGTATTCCTAATACCTAAGCACTGTCCCAC
ACATGGTAGATGTTCAATACATATTTGTTGGATAAATAAAAAATACAGAGCTCCCTCTTCTCCCCTACATGTTGGAAAA
TGTGTGTGATCTAAATGAGGGAGAGGGAGAACCATCTCATCTAATCTTCCAAACTTCCTGAGCCACCTTTAGCTACTGA
GCACATTCACGTTTCCGTCAGCTGGGAATTTAGGTGCCCTGCCTATCATCCTCTGGAGCCCATCCCAGGAGGGGGCAGA
AGGAGGTGGAGCAGCACACTAGTGTTCCCAGAGTGGGGAGAGCAGATGGAGAGGGAGTTCATGGGAACGATGATTTACCC
CTGCTCACTTACCCCTGTGAGAGGACACTGCCCTCCACGAATCAATGCTGGCTCCTCCTCCAGGAACCAGGCTTCTGCA
TCTCACTTGCCATGCTGTGGATCTAGGAACAAATTTGCATTTGGAAATTCATTTTAAAGTGTGCATGCTGAGGTTCTGG
TGACATGCAGGGTAACTTCTCCCACCCACGATATCTCTTGACTCAGGCCTTCATGTAGGAGGGGAGGAGTGTTTGGCAG
CAAAACTGGGTATTTCTTAGGAAGGAAATCCTTCTAAGCTTTATGGAAATCCTAGGGTGTTGTCCCTACCTTTCTGAGT
GTGGAGCAGACTATCTGTGTACATAGCGGTAGTGAGGGGTGTAATATTGGAGAATGGCTTAGGTAACAGCCAAAATATA
CATTGTATATCCCCAAAAAGACGACCAAAGACTCACAGGGCCATGCTGCTATCATTAGATATAATAGGCAWAGCCTTTC
ATTGATATGTAAATGTTAGTATTTTTGGTAAGTGCCTTATTTTATCTTCTGGATAATATAAGATCAAAGTACCATGAGG
GCAGGATCCTTCTCTCATTCATTTTTTTTCCCAAGAGCATAGCACAAAGTAGGAAGTACTTGTTGAGTGAATGAATGAT
CCTAAATATGAAAGGATGGCATTTTATCGCCATTTTCCAGATGAGAAAAAAACGAGGGAGCTTATCAGGTTTATATATA
GTGAGTGACAAACTTCAGAATTAAACCAAAATCTAGTGATTCCTATAACTTACAGGTTATTAAGGAGTCCACCTGAATG
TAAGTCCAACAAGACAGGTGTTTATTTTGTTTACATACATTCCAAGTACCTAAGACAATGCCTAACATATATAAAATAC
TTCATAAACATCTTCATGTGTGAATAGAATTTTCAAAGTTCATCTCAATATATGCCTCATGTGACACTTCCAACCCTTC
ACTGCCAGGTCTTTCAGTCTTTATTTGAATAGGTTCAGTAAGAAAGAATTCACTCGTGCTTTAGATATGTATGGCTTTT
CCACATTGTTGTCAGTTCTAAATCCCGAAAGTTATTTTGTATGCTTGGAATACTTCACAATATTCACAATTTTTACAA
TAAAATAATTTCAGGTCTTCTTGATGAAAAATGTCGATCAGGACAGGGCTGAGGATAATCACAAGGCTTGAAATTAAAG
AAGTCTCCTTAGGCTGAAATGAATTTCTAGTTGTAAAACCAAATATGAATCTATGTATTTGTGTTCCTCAGACCACTTT
TCCAGCAGGTCCTTGGCTGTATTTTGTATGCATTTGTGAAATGCTTTGTTGAAATTCAGTTTCAGCCAGGTCTGTAGCA
TTCTCCTGGCCCCTGGTCTGATAGTTGTACCAGGAACACTAGCTAACAATTGCATGGCACTAGTCTCAGCTGGATTTGT
TCTAACTCTTTTATATGTATTAACTCAATCCCTACAACCACCCCATTTCATTACTATGCCCACTTGAGAACTRAAAACA
CTGAGGAAGAGCGAAGTTAAGTAACTTGCAGAGGTCTCAGAGCTAGGAAGTAAGACAGCCAACTCGGAACCCAGCAGAC
TAGCTCTGGAGCCTACCCACCTACCCTCCACACCGTACCTGTGGTCCATCCTCACCAACCAGGCCTGCTGCCCTGAGCA
TAACTTCACCCTCTTGCGTTCTAAAACTATCTGCTGATTAAGTGAGTTCTGAAATGTTTTCAGGGACCAATACTAAAAT
CAACTATGAAATAGTTTCAGAGATTGACCTTTCCTCCTTTTGGAATGGATAAGATTTTCCCAACCGAGGTCTCCCATCA

Fig. 6.172

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GCTCTCCTGTTCCCCTGGACTGTGGCACATAGGAGGGCCAGGGACTTGCGCCTATTATTTCATTTCTTTTGTTATCCTT
CATTTCTAGATCTCTTCTCTACCACAATTATCTCTCATGTTGACAGTTTCCCCACAGATGGGATGTCTATTTCCTTGAT
AATACAGCATTATTATGCTAAAATCTCTCATTTGTGTTATTAGCCAGACTCTGTGCAAAAACGTGTGTATATCTGAGAC
TCTGTGCTGTTCTTAGCCACCTTCTTGGTCCCTTCTCATGTAAATTACTAGGGCACTTCTTCAATATTATTATTATCAT
GCCCCCTTACCACCACCACAAAATCTTCCATCATAACCCATTTTTTTTTCCTCCTCCTTACCACATTTTCAGCTTTGAA
CTCTTATAATCAGATCAGGAATTACATTTCTGAACAAATGCAGGCTTTGCAGCATTGTGAAGTGCAGCTGGTCCCCAGC
AGAGAGCCCTAAGCCCCTGCCTGAAGCCAAACCCCCTTTTGTGGCGAGAGGTCTGGATTTTATTTGTTGTTTTTATCTT
TGGTGAACTGTCAACTGGAAGCAGAGATGCGAACACCAGTTATGTCCTCCCCTGCAAGGTTCATGACCAATACTTTATA
GTTTCTGGATATGCTTCCTAGGTTTCTTCTGTTGGTGTCATTTGCCTGCATGTGACTCACTGGCAGGTGGTGATTTCAT
CATATTCTAGACATGTGTCTTAGGAAGCTGGCCCCTATCCTGGCTGGCTGGACACATATCGAAGTTCCTTAGATCTACC
ATTGTCACTTTCATATCCATTGTTAAATACTATAGATAGCATGCAGGGGTCAGCCAAATTGTTCTGTAAAAGGCTCAAG
CATAAATATTTTAAGTTTTGTGGGACAGACAGTCTCTGTCGCAACTACTTGACTGATGCAGCATGCAAGCAGCCACAGA
CAATAAGTAAACAAATGGGAATGGTTGTGTCCCAATAAAGCTTTATTTATAAAAACTGTCTTTTTTTTATAAATAAAGGA
GCCCATCGTTTGCCGCCCCCTAGTGCCAGGAAGGAACAAGGGGAAAATAAGGAAGAGACCCAGAGGAAAAAAAGTGTCA
CCAACCATATCTCTAACTCATATTTATTTACTTATTTTTCTAAGCGCAATCGCTAAATTATCTCGTCTTCCTAAGGCGT
ATGAATCTCTTACATTAACTTAGGGACAGAATATAGAGTTTTATGTTATCTTCTGTGAGTTCTTTTTCCATTCTGTCAT
GTTTCAGACTTTCAGAGGTAATTTTTAAGTATTTCAGCTGTGCCTTAGATACATTAAGGGACCTATAGTGCCTCCCTGA
GGCCAAGATGATGGTTGCAATCATAAATAGGTTATTTCTGAGTTGAGACTGGTAAGAACAGATTATCCTGCCCAGACAG
TGATGTGCCAAGCATTGGCAACTTTGAAAAATTTGCAATTTGATTTTATAGGAGAGTTTCCTTAACAGGCCTGAGTTTT
TTTCCCATTGCTTTTGTGAATACAGTATTTTCCCCTGCTAGACTCTTAGTCAAATGTTCTTGCCATTATAAGAAAAAAG
GAAAACTGAAAAACTTGAAAATAATTTTTATTGGCTTAGGAATTATTTTCATGTTTGTCTGCTAATGGCAAATAAATCTA
CCTTATTGAATCATACGGATTTTAATTCCCCGGGGGAAAGGTAACCAATGACAGGCATATGTGTGTGTGTGTGTGTGTG
TGTGTGTGTACAAGTGTATGCTTGTGAAAATTGAACAGCAAAAAGAAATTAGGTGGAGAGGAAAACATTAAAGCTGG
TACTGGTGGGTAATCATTATATATAGTATATATACACATATGCCTTCTCCAGTTTTATGGAGGCAGTCAGTGCCTCAAA
GACGTGTAATAAAGAGACAGCAACAATTATTGGTGGCATTCATGTAGGAGAGGCTTGAGCCAATGTAAACTTGACTTGG
TTAGTAATCTTTGCCAAAACTCCCTGATTGTYGTAAGGGAGTTGTGAACCATTAGGTTTGCTTAACCAGAACAGCAAGT
AAGCATCCAAACAGATTGTCCTTGGACCATAAGCAAATGCCCAGAAAATAGCCCCTGCATAGGAKATTAAGTACTTAG
AAACAAATGAGAAAAAATATTACAGCAAGTCTTCCAGGTATAGAAGGGATAAACAACCCTATTGGTATCAGAATTTAGG
AATATTATTATGTCTCCCACTTTAAATGAAATACAGCAGCCTGACTTGGCAAAATTTGGGGAACGGTTCCCCTTCTTTG
CAGCCAGGTTCGGTGCTTGCCCTCTGGCATCCCTTCAATTGTGTATATTACTGGCAGAGACAGCTGTTTGAGCTGCTAA
TAAGATTGCAAAATAATATCAGCTTGTGCTTCAGGTTGAAGATAAGAGATAACATTGCATTGGCTGTAGGAATGGCTTT
CTTATAAACTGGATTTAGTTTTATCTCACTTGTCTCACTTGTTACTTTTTTATTTATTTACTTTTTTTTTTTTGAGATG
GAGTCTCACTTTGTCACCTGGGCTGGAGTACAGTGACACGATCTTGGCTCACTGCAACGTCCGCCTCCCAGGTTCAAGC
CATTCTCCTGCCTCAGCCTCCCAAGTAGCTGGGATTACAGGCATGCGCCACCACGCCTGGCTAATTTCTGTATTTTTAG
TAGAGACAGGGTTTCACCATGTTGGCCAGGCTGGTCTCAAACTCCTGACCTCAGGTGATCCGTCTGCCTCGGCCTCCCA
AAGTGCTGGGATTTCAGATTACAGGTGCCATGACGCCTGGCCCCACTTGATACATTTGAGGGTCTCATAAGGTCTTGAA
ATTGAGCCTGTGAATTTATAAACAGGGCTACAACCATCCTGCAGATGCAACTTTTTGGCATGCCAGAATCAGTGGCACC
TGAAGAGCATCTCCTTGAAAGCGGTAGTAGATAAAAGGAGATGCCTTTCAGGTTTCAGACGTGGACCTCAGATTCACCT
CCAAGACTAGAGTTATCACAGTGCACAGGATCATCCAAGAACAGACATGCAGAGGGAGGAAAGGCCATTGTAAGGCTCA
GGAGAACATAGTAGAGCCTAAAAATGAGAACAAATGGTACTATCAGAAAAGATGGCAGTGGTTGGGACATAAGAGAGG
GTAGGTTCACAGGTTTGTGGGACCAGGACTCAGTCACTTTCCTCGCGCTAGGCTCAACCCTCACGATGGGGAAAGTAAT
GTACTTCTGGGCTTAAAAATTCTTCACAGTCAACTGGTAGAGATATGAGGGAATTTATTGCTAAGTAGTAGGGAAAGTGT
TTTCACAAATAACATACAGGAAATTCCCATATCATACAAAGGAGGTTGTATTATCACCACGTTTTTACATATGAGAAAA
TAGAGTCTTAGAAAAGTTATGTACCTTGCCTAGGAGCATGTGCCTAGATAGTGGCAGCTCCAAGATTCAAACCTAAAAC
TCCAGAGTCTGAAACTCACACAGTTTCTTCTCTGTTCTGAGGCTTCATGCAGCTGGACAGGAGGTTAGCCTGAGACAAG
AGGAGGATCATTCTCGAAGGCAGAGAGGCTGGGACATGACTGCTTTCTGGAAACCTGAGCTCATGGAGAGAGGCTGAG
AGTCAGGAAAGTAGGACATGAAATCATTTAATGAGAGTGAAGACCAGCCAGGTCAGGGAGTTTGAAAAGAGAGTTTTGA
AATATCCATTGTAGCTAATATGGGAGAGCATTCTTTGAGAGTGATTAAAATGGCGACTGAGGACACAGCTAAAGTGAGA
AAGAATAAATCTGTGGCAGACTTGCATATTTTTGGTATGTGCATCTCTTTCCGAAATCAATTGAACAGAGGTGGGAAAA
GTAGCTTGGTGGATAATGAGGTAGAGTGGAAAGTTGAAAAGACAGAGCTTAGGAGTCTTTCTGATTGGTCTAAAATTCC
CCAGGGAAAATCATGTTTCTTTTTATGAATATAGTCCAGGTCTTAGGTCTGTCCCACGGCACAAAGTTTTCAGGACATT
CTGGACAAATTCACACATTTCCCTGGATGTCAGTCAATTCCCTTTGCCTCTGTGCTATCACTTTCCTTCTCTCTGCTTT
CTCCTTGAGTTACCTGTCCCAGCTCTTCACTCCACATTCTGCAGCCTCCGTTTAGCTCCTTTATACCTTTGTCCAGATT
CTTCCTAACTTATTATACTGCCTCCTCATGTCAACTTCCATCCCATCTGCATTTGGGAAAAGAAGCCCACATGATAAGT
GAGTCCATCTGCTAAATGTATGCATCTTAGAGCATTTGTTCAAGATCAATAAAGATACAAATTATTTTGATTTTCAGC
TGTTGCCCAAATCCACAAGAGTACACAAATGAAAATGCACTGATTTTTAAAAAGTACAAATAGCCATTAAAGTTTGTAT
TGTTTTATTCTTTAAATAAGCCGGAAAAATGGGAATACGTAGGAACGAGGGAGCTAGCAAAATAGGCGGGGCTTCTGCG
TTTCAGGTGGTCTGAGACTGCAGTGACCCTGAGGCTGGTGGACCTGGAAGGAGCCAGGGAGAGATGGTGGATCCCAGGG
GTGGGAGCAAATGCATGGTCCAAGGTTCTGAATGGCAGAGCTAGACCAAAGGACTAACCTGCTAACCTTGACCAAGGAG
ATGGCCCAGAGGTTTTCTATTTCTACTGCTCCATCTTTTCATTATAAGTTCCCAGTAGAAACGAGTCATCATTTAAAAT
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Fig. 6.173

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ATCGGTGCAGCACTTAACACATAAGGAAAAATCAATAACATTCATTTACTTACAGTACCCAGAGAAGAAATAAATTCAA
ATACGTAAAAATGGGCATGAACCATYCTGTTCCATATCCCACTTTAGGACTGTCAGCAACTGTATACATTGCTGAACGT
CATTGCTCTTGTACATAATGAATAATTTTCTGTAACCCTGTGAAGGCCGACTGATTTTTACGGAAAGACAGCTTTGTGT
TAGGCGGCCCGCCTTCCCCACGGAATCGGGTTTTCCCACGCCACGTTTTGATGTTTCTGAGAAGTGTGCGCCATCTGCT
GGCCGCTGAGAGGATTGACGAGCAGCCATAAGGAGCACCGTGTTTCTTGAGCATGACTTAACCAGGCAATGGAATTACA
ATTATCTTTGGGTTTCTAAAACATAATAGACATTTCAGGATTCACGTGACTTATGAGAAGTCGCTTAATTGCCTTTCAA
GCAGTATACATTTTTTATTCAATATGTGTATTATTCTTTGTAATAATAAACRATGCAATAAAGCAATCGAGGGGTTCCC
ACATGCTCTCTAGGCAGAGATGGACCCCGGGCCCTAGATGAAAGGTTAATAGTTTGGAGTGAGCACTAACTCTGGTYCT
CTGACTCTGAATCTAAGTCAGGAGTAGAATAGAGCTGTACAAAGAAGACTGCTTGATGGAATTGAATTTTGTCCCTTCT
TCAATGGGGGAAAATAAAGTACAGTTTTCATCAATAGATTAATGACTAAACTTGGACCTTGATGCCAAATCCAACTTGA
GCTCCGCCTGCACAGTCTGGAGATGTACATAAGAGACACATGCTCAGCCTCTTAGCACATCGTTATAATCCAGGCAGGC
AATACTATATCCTAAAATTACTCCTCTTTGCTGAAATTGAGAAAATAGGAATTACCCAGGGGTAAATTTGGTACATAAA
TGTACCAAATGGTATGATTTATTCTAGTACTGTCTTCTTTTTCTTGATTTTCTTCTTTATTGGGAAATTTGTATTTTGT
AGTTCCTTCGACAGTTTAAAAATTAAACTAGAGGGTAAGATTTAAGTTTTCTCACAACTAACAAATGATTTTGATTTTA
TTGCCCAAAGCCCTTTATATCTCACTTTCGATGGGTCTAGTCCCTTTGGCTTAATTTAGATGTGATTTTTCCTTAATAA
TTTTGAAAATGGTGGTGCTCTATACCAACTAGCTAGAATAGGAATCACATTGACAGTATTTAATGGATAGAAGCATCTC
AAAGATCTTTGCTTAATGAGGGCAAATCATTAGCAATGTCCAACTCCCTATGTTAGAGGCGCCAACTAAGGCACGGTAC
TTAAATTTAAAGCACAACTCTTCGACTTCCATAAAGTACATATATTACATTATATTTTTAAATTACGGGGTAGTGACTC
GATATGTCACTGGAATGTTTTGAATAGTGCCCCCGGCCCCCCTGCCAAAAAAAGAATGCAAGAACACTGTCCTAACCA
AGAAGGCCTGGGAGTTTTAACTAATTCAGCCTTTCAGCCTCCACTTAAAATTCTGGAAGACAGTAGTGTTGGTGCTAGC
TGTCCTTGGCTGCAAGTTGTCTTTACAGAAGAACTTGGGGTACATTTTGGTGTGAGGCCACTCTTGCAAGCAGCATTTG
CAGGCATTAAGAGAATGCACACAGTTCAATAAGCTGGGGGTTATATTCATAATCAGTCTTTGTTAATGCTGTAAAAATA
CTAAAGGAATGGCCAAAATCTCTCGGTATCTGGGGAAAAGGGTTCTTTGAAACTTTTGATGTGTGAGCAGAATAATGGA
AACTTATCTGTTCCCCATGCTTTTAACACACACACACAAACACACACACACACACAAACGTCATAGATCAAGATGTA
ACAGGTTGGAATGGGGCCTAGGCTTGGGAATTTTGTGACTCCCCAGGTGATCTCAATGTACTGCTGGTTTATTAGCAGG
CTTTGCCCAAACAAATACTCTTTCTCACCTACCAGGAGAAATCAAAGAGCTCTGGGTTAAACACTTGTATGTTAAAACC
AAAATGGTGGGCATTGCAGAGTAGAGGAATTAGGAGAAGAAAATAGATATTGGGTCAAAAATTGGCCTCAGGAACCAGG
AAGTAAGAATGGCCAATCTTAAAAGAGAAAGTTCTGTTGAAGAGGGAGCATTATCTGAAGTTGGAATGACTATGGAGGA
CAATGACAGCTGTCCCACAGGATGTCACTGGGCTGAGGAAAATCAGACTGGAAGCCACAAGCCCATGGGGGGCTCAGGAA
GCCTGTTTCCTGAGCAGAACAGAGCTGGGTTTGGCACATGCCTAGTGTTCCAGTGAAATCTGTCATTTTTCTGCTTCCA
CTTTTTACAAATGAGTTATTCAACGCTACTTTGTTGACACATGGACTCAGCATTTAAAAAATGACAGTGGAGATAAGGG
TCTAAGGTAGTGATAGGCAAATGAATGATGGGCATCCATCATCTGAAGAGCTTCTTAAAAATTCAGATTTCCCAACCCCA
CCACAGACGTCTGTCTCCGTAGGTCTTGGGTAGGGCTCAGAGCAGGCTGCTTCTGAAATTTTGAGGTTGTGAGGTAAGA
ATACAGGTGGGAGCCTGCATACCACTTGTCTAAATATTTAAAAGTTGTAAATCAGCCTAATGAACTGTGAAGGAAGCTC
TGGCCTACTTCCTTGACAACTTAAGCCTATACAGAGACCTGGAAGATCAAGTTCAAATGCAGAACCTGAATGTTTCGGT
TCTATCTGCTCCCTCAGTGTTAAAGGTCTGTCCTTTTTCCTTGACAAATATGCTTTCATAATTGTCTGGAAGGCCGGAT
TCCAATTCAGCAATTCCAATCCCTCAGTGTTGAGGGAAAGGGCCACCCGCTCTGCTCTTCTCTTCCCATCTGGCTCTGT
CTGGAACCACGAGCACCTCTGTGTGTGTGTGGGGACAAGCCAGCCCAGGTGCCTGCGTTTTGGCCCAGTACCTCCTAC
CCTCATACTACATATGTAGCCACCCCCAGGCATAGGGTTGTGCACACTGGTGACATGTCTGCCCTCTAGAAGAAGCATG
GGGGAAATGGGACCCCACAGGCCCTGTTAGCTGATACGGGCCATGAGCTCCCAGGTCACGGTCTAAAAGTACATTGGTT
TGGGGCTCTGCGTGACCATATTCCAAGGATCCTGCAAAGGAAGAGGCAGGACTGGAGAAGGGAAAGAGTGGGCCTTCTA
AGGCTCCCAGGGCAGGACCTCTTTTTGTCTGGTCTAAGAGGGGTATTGAACCCAAGTGGCTTAGATGCTGAATCCCTGG
GCTTAGAACCCAATGCCCTGGAGCATTGATCTCACAGTGCTGGCCAGAAAGGGGCTCGTTGGTTGCTACTAGGTCATTA
TCACTCCCAGGGCACCTGGGCAGTCAATGAAACTGAGCTCAGAAAGTCCTGCCTTTATGATGTCCCTTCCACTTCCAT
AACTTTCTCCATAAAATAAAACCAAGAAAAATAATTTTTCTCTTGATYTCAAATCGTATTTAAAGGAAAATAGAAATTT
TATTTTTAAAGAAATTGCATGTTTTCATTGTCTATATATTGTTTTGAAATATGTACACGTTGTGTAATGGCTAAATTGA
GCTAATTAAATGTATAACCTCACATGCTTATCATTTTTGTGGTGAAAACACTTAAAATCCACTCTCTTAACAATTTTCA
AAAATATTTGAAAAAATATAGTTATTAACTGTAGTCCTGTAACTATTTCCTATTAATTATAGGGAAGTAGAAATTTTAA
TAGTCATCTTTCTCATGCTTTTCTTGACTTTTAGATTCAAGAAATAATAAACTTGAATAAAAGAGATTAATTTCACAGT
AATGCAATATTTCTGTAAAATTTCACACCTTGTTGACTGTCAATCAGCTGAGAACTTTACACCTCATTAAAAATCCATT
AGAACAATGGCTGTTCCATTAGCCCTTCAGTTTATGCCTTAAAACATACTTGTCTAAAARCAACATTGTTTTTGGCAGG
CTGCTCCTTTTGATAAGTGGGGACTTGCTCTGGTATTAACTGTTTCCAGGTCTCCAGAAAAAGGCTCAGGGTATGTCTA
GGGACATCAGTGACAACAGTGAGGTAGAGCAAGTCAATACTGTGCCACCTCTGCGCTAGCCAGAGAACCTCTGCTCCAA
GGTGAGTTGGTTCCCCCTTTCATTAAGGTCCTCAGGCCCCAGTAAAATCCAGATATGGCTTTGGAGAGGAACGCTGTCT
GAGTGAAGCTCCCTCAGTTTCATAACCAGCTTAGAGCTTCTCAGCTCTGACCTGCCTGTTAGCATGGCTGGGGAGCCCA
GTGGTTTATTGAACTGGCTAATAACAAAGATTCTGAGATAACGAAACAGAAAACAGGTAGGTTTTGATGAAAAGGAAGG
AGATAAGGAGGATAAGAGGGGTAGTAGGGAGGAATAGTGGGAGAGGGGCAAAGAAGCATTAAATCAAGATTTGGAGAAT
GGCTGGAAACAGTGGCAGAGACTCAATAGGAGAGTGACCCTGACAGACGGGGAGGTGGCAAAAGTGCTTGCAGGCACAG
GCCACCCAACAGTTATTTCRGGGACTGTTCACAATCTCATCTACTTCACAGTTTTGCCTTGTCCTAACTCAGGTCTTAT
TAGAGCCAGAATAAAATTGCCTTGCATCGTCAGAAACTGACCATTCATTCCTGTGGTCTTCTTTGAGATTTTTAGAATCC
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Fig. 6 (174)

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CAAGGACTAAATATCCCTATCCACTTGCCTTTGTTCACTGTATTGGATGATGTGCCATTGCCTGTCTCCAGCATTCACT
GAGAGCAGTACCCCAACATCCAACACATCTTTAAATGCCAAAGGGAAGTGAAAATGTCCTGACTTCTTAAACTCTAACA
TAGCATTCATCCTATTCTTATATTTTTTGAACTTTGCATCAAAACATACATAAATGCAGGTGCACAAATTGTAAGTGTA
AAAGTCAATGAATTTTCACTAAGTGAACACACTTGGCTCACCAGAAGCTAAAGAAAGGGTGAACTAGAAGGGGACTTTG
TAGATTACACTTCCTGACACCATCATTATTTAAGAAAGAACTGTTTCCTAGGTATATTTTAGGACCTTTTATCTGGTGA
TATCTATCTACTAACTTTAGGGGTGGGGTGGAATTGATTCTTTTTACAAGATTAGTGAAGCTAGAAACAACTAGCTTTT
AATTCATTAAATGTAACCTACTTTCAAGTAGCTTCATGTGACTGAAATCAATCTGTTTTATTTTCCAAACCAGTGGAAT
ACACACAGGGTCATTACATACATTTTGAAAGTCTTCTGTTTCAAATGTATTGTAGATTAAGAAACAGAATGTGGCAGGG
CACGGTGGCTCATGCCTGTAATCTCAGCACTTTGGGAGACCGAGGCGGGTGGATCACTTGAGGTCAGGAATTTGAGATC
AGCCTGGCCAACATGGTGAAACCCCGTCTCTACTAAAAATACAAAAAATTAGCCAGGCATGGTGGTGTGCGCCTGTAGT
CCCAGCTACTTGGGAAGCTGAGGTGGGAGAATCGCTTGAACCTGGGAGGCAGAGGTTGCAGTGAGCTGAGATCATGTCA
TTGCACTCCAGCCTGGGTGACAGAGCGAGATTTTGTCTCAAAAGAAAAAACAAAAAATAGAAAAAGAAACAACGCTATT
GGCACACCCAGAATAATCACTATTGGACTTCTGATACCAAAAACAAGGGGGATTTCCTCGTTTTTAAGTTTTATGAAAT
GAAGTAATATTCTGTATACTCTTCCGAGTTTAGCTTCTTTGAGTAGGTTTGTGAGATTTATCCAAGTGTGGAATGGAGT
TATAATTTATTCATTTTTGGTTTTGGGTGATTGTATTTTTAAAATTAAGGTAAACATTAAAAGTTAGGGTTTAACCCAC
TACTGGGTATTCACCCAAAGGGAAAGAAGTCATTCTACGAGAAAGACCACATGCACAGGCCTGTCCAATTCACAGTTGC
AAAAATAGGCAACCAAACTTGTTGCCCATCGACCAACGAGTGGATAAAGAAAATGTGGTATATATGCCCTATGGATTAC
TACTCAGTCATAAAAAGAAACCAAATAATGTCTCTGGCAGCAACTTGGCTGGAAGTGGAGGCCATTTATTCTAAGAGAC
GTAACTCAGGAATGGAAAACCAAATATCATGTTTTCACTTACAAGTGGGAGCTAAGCGCTGAGGATACAAAGGCATAAG
AATGAATAATGGACTTTGGGGATGGGGGAGAGATTGGGAGAAGTTGAGAAATAAAAGACTACATATTGGGTACAGTGTA
CACTGCTCCGGTGATGGATGCACCAAAATATCAGAAATCACCACTAATGAATGTGTCCAAGTAAACAAAACCCACTGTA
CCCCAAAAACTATAGAAATAGAAAATGAATAAATAAAAGTTAGGGATTAAAAAGAAAGTGGGACATGGTCATTCCAGAA
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TCCTGTGGAATGGTTCCTAGCCCACGTGACCTTGTCTGTGTTGAAGCATGAGTGAATACATACTACCATCTACTTTAGT
GAATGTTGATGATAAAATGGCCAATTTCATGTATTTGTTTATTTTCAAGAAAACCTTAGAGCACTCGTTCTCAAACTTT
AGGATGCATCAGAATTACCTGGAGTGCCAGAATCCCCTGAAGTGGTTGTAAGAACTCATTGCTGGGGCCAGGTGCGGTG
GCTCACACCTGTAATCCCAGCACTTTCGGAGGCCAAGGCAGGCGGATCACTTGAGGTCAGGAATTCGAGACCAGCCTGG
CCACCATGGTGAAACCCCTTCTCTACTAAAAATACAAAGATTAGCTGTGCGTGATGGCGGGCACCTGTAATCCCAGCCA
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Fig. 6.175

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AAAAAATTAAAAAAATGTTTAGCACCTTCAAATATCAGATTCACATCCTCATCATCCTCACTTAGAGATTTTAAAACTAA
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Fig. 6. 176.

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Fig. 6.177

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TAGAGGAAGAAATCAAGGTTATTCAAGTAGAGGAATGACACGTTTTAGAAAGACTTTGTTATAACGTTATGGGAAATG
GATTATAGGCTGGAGGCAAGAGATAAGACATTGTCAGTGTGACAGAATATTTTATTAATAAAGCACATAATTTATACTC
AGATCTATGTGAGATGAGACCAATTCATGTCAACATGTCCAATGTTTTTGAAAAAATGCTAAGTTTTTATTATCGCTTT
TGTTTATTTCCCTGTGAAATATAGTAGGCTAGCCCTCTATTTCTCCTGGCGAAATCTTTAGATAGGTGGAAAAAAAAAT
GTTGAAAAGGGCATCAATGACCTCTTTTATCTTGGTTTTATCTGCCTTTTGGAAGATACTTGATTCAAAGCTATGAAA

Fig. 6.178.

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AAGCATCTCACCAATCCCGTATTTCACTTGGTTGGACGATGTGTCAAGAGAACGCCTGAGGCTTCCTCTGTCCGTTAAC
AGGCCAGTCATGAGTCACAGTCTGAACATGAGGAAATGTTCATGTACTTTTTCATAGTCATGGCAATGAGACTTTTTTG
TGCCCAAAGTACTATCCTCGCCTAAAGTCCCTGTTAAAAAGGTTATTTTCTGCCACAGGCAGGTAGCAGAATGTCAAGA
TCCAATTCCAAGTGCTTCATCATCACTCGGATAGCCTCAGGAAGCTGAGGTTTTACAGCTGCCTGGAACCCACATGCTT
CATGTGATGTAGAGCAGCCCTGTTAATTAGCTGTTTCAGGATTCTTAATGCCCTATGGCTTTGGAAACACTTGTTAGCC
CACAGAGGGAGGGGAAGAGCTGGAAGCAATTGGACTACAGCCTGAATTTATTTTAATTCAATTCAGTAACCACCTTATC
TTTCTAAATCCCTGTTTAGAATCTGTTGAAAATTATAGTTGATTTAAGTTTTTCTCAGTGATGCATCCTGAGAGAAGAG
GGAGATTACAATTCAAGGTTTTCTGAAGCCACAGAAATGTGGAGTGCTGCAGAAAGAGCTGATCGATGGAACTTTCCAG
ACACATGTATGCAATTTTCTAAAGGCTAGAAAATAGATGCCGGGGATGGGGAGAGGGGAGGCCAGTTGATGTGGAAAGG
CTTGTGTAGAGTGCCCACGAGTCTGCATTTCTGTTCACCTACCAGAAGAAACACTGCCAGTTCCCCCCACCCTCACCCC
CCAACCATTCTCACTGCAGAGAGCTGGCACTCAACAAACTCATAATTTTTCTCAATACAATTCTCATGACATAACCTGA
CCAACCATATGAAACCTAATGAAGAAAACAACCATTGCCATTGGCAACAAATGCAGGAGTTGCATCACAAGTAAACCAC
ATACTTTACTTCACAAATTTGAACATTAGAGTACCACCGGCAAAAACATGGTACCATCCTTCGGTATGGATGAGATATC
AGTTTACAAAACCTTCACTGGAAAGTCCATACCAAAATTTTTAAAATGCATGCAAAAGCTAATTGTGTTGACATCTTTC
TTATCAGAGCTATTGCAAATTATTTGTCTAGAGGTGGTTCTTAAAAACAGCCACCTGTCTGTCCACTAATCATGTACAT
AGCATCCTCTTCCTCATATTCATCCTCTGAAATACCAAGAGGTGAAAGAGTGTTTCAGTAGCTCACCATTCTTGGCTGT
TACATAATGGGTCAGATTCCTTTCTGAGTAGTAGGAGAGGTCAGTGATCCAGATCTATATATCTTTGTGCTTTTTCCAC
TAATACATTCTTAGAACCTAGTAGGTGCTCAATAAACGTTAGCTGTATGAATGAGTTAGTGAATCAATGAATGAAGTGT
TAATTTTTAGAAATATTATGGTCAGAATTTTACTAGGAAAAACATTTCTTATAATCACAAATGAAAGCTAGGTTTACTC
AAAATTGGGAATAGTAAGTAGTCTTTCCATTTCCCATACCACAGGTATGCTTGTATTCTTCTTATATGCTTATGAAAGA
AGAGAATGTGGATTAAGTTATAAACTATTTGTACATGATATTGTATATGTAAAAATCATGTCAGTTTTGAGGCCAGGAT
TTCTTTGTATTGAGTAAAATTTTCCTAAATATTTGCCCATTGTATTAGTTCATTCTCACATTGCTGATAAAGACATACT
CAAGACTGGGTAATTTATAAAGAAAAACAGGTTTAAGGGACTTACAGTTCCACGTGACTGGGGAGGCCTCACAATCATG
GTGGAAGGCAAAAGGCATGTCTTACATGGTGGCAGACATGAGAGAAAATGAGAGAAAAACCATGAAAAAGGGGTTTCCC
CTTATAAAGCCATCAGATCTTGTGAGACTTATTCACTACCATGAGAACAGTATGGGAAAATTGCCCCCATGATTCAAT
TATCTCCCACCAGGTCCCTCCCACAACACATGGGAATCATGAGAGCTACAGTTCAAGATGAGATTTGGGTGGGGACACA
GCCAAACCACATCACCCATTTTAAAAACAGGCTTGATATCAATTTATTGCTAGAAAACTATAATTGTATTTTCTTTTAC
ATTTTTAATGACTGATTGTACATTTGTTCCCTCAAAAGAGGCTCTCCAAAATACTGCCTAATTCCTATGTTAATAGCAA
CCCAAACATGTTTTCAATACTAATATTAAAACAAATTAGCTTAACAAATACAAAAGGTAATGCCTGGTTTATACAATAC
GGATTTGAGAAATGCAGGTTTCAACCTGTTCCACACCATGCCCTTCATTATATTCTTCTGCCAGAGATTTTATTCTTTA
CTTTTCATTCAATCAGAAAGAATTGAGAGGAATTTGAACCCATTCTGCCTATTTAGAATATCATTTGTACTGACTTTAA
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GGCCCAATTACCAACTCACTGCAGCCTTGACCTTCCAGGCTGAAACCATCTTCCCACCTCAGCCTCCCAAATAGCTAAG
ACCACAGGTGCACACCACCATACCCACTAATTTTTTGTAGAGATAGGGTTTCGCCATATTGCCCAGGCTGGTCTTGAAC
TCCTGGTCTCAACCAATTCACCTGCCTTGGCCTCCCAAAGTGCTGGGGTTACAGGTGTGAGCCACCATGCCTGGCCAAG
TTGCCTGGTTCTTTACATGGAATGTTCTTTACATGGAATGTTCCAGGATCAATTAAACACAGATAAATAACAACTTATG
TTTTGTAAATAAACACATCTGAATGGAATCTTACTCCAATTTAGATAGAAATTGATATTTTTCCTTTCAAATGAAATAA
ATGTTTTAGACTAGGACATCCCAAGTGAATTGAAGATTGGTACAGGAGCCTGAGGAAGGAAGTTTAAGAGAGTGAAAGC
ATGAGAGAAGAGCCAAGATTATGTAGCCCAGAGAAGGAAAGAATTATTACCTACCATTACTTTAAGGGACTGTGCTTCC
TCACAGTAGTGGAAGACAGATGCTCATTTTTTATTCAAGACTGTGTAAAATAAATGGCCTTAATTTATAGATGTGGATG
CTGGTTTATATCAATACACCTCAACCTGGGATGGCTCCAAATCTTGACTACACATGGGAATAACTTGAGGAGTTTTACA
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GATGCAAAGGGAAGTTGGAGGAGGAAGAGAAGGAAATGGTAAAGTCCAAAGCATGGGTGAAGGGGGGCCAATATAAACC
CACAGAGCAAAAAGGAGATGTTGGGTCATCATATTCTATTACATAAATTGGATTTACATTTCCTTGTCTTTCATTGCTG
GTCTTTATTCCCACAAGTCTCCAAATACAGTCATAAAATTCTCATGAGTTGTTATAGCAAAATTGCTTATCATTTACTA
TTTCTTAAATGAATGCATAAACTGTATTACTTTGGCAGAAAGGATGCTGCTGGGTATCATATGTAATGTATACTAGTAA
GGTGGACAGGACCATTAGGACTTTAAATTCCTTTATAATCTCAAAAGTCTGTGATTCTGATCTTCCTGGTCTTGCAACT
CCAGAAAAGGTGGTAATCACTGGAGTAGGCTATTTATGGGCCTGTGAGATAGTGAAACATGCTATTAAGACAAAATGAG
AGACTTCCTTCTCTGAAATGGTTCCATATAAAAGTAATATATGGCTTAAGTATCCAGGGGCTCATGTAATCTCTCCATG
TCAGTATCTTTTACGGGGAATTATTTAACCAAACATGTATTGAACACTTACGTGCTACACTAAAATCCCAGATTTGTCA
AGTTCCCGTTTAAAAAGTGGGTACTAAGCCTAAGATATTAGCAATACTTGTATCTGAAAAGGCCTCACATCTAGAAACT
ATAAAGAACTTCTAAAAATCAATCAAAACAACATAGTTCTTAGAAAATCACAAAGGAACAGGTTTTTCAGACTCGAGGA
TGTCTATATGGTGAAAAAGCAGCAGTATTTTAGAGTTGGTCTGAACTGACTCACAGGAACCTATTTTTAAATGTCCAGA
AATTTGCTAAGCAAATTGTTAATGCATCAATTATCTAAAGTTTAATTATGTAAACTTCAAGTTATCTGAAAAACATAAC
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Fig. 6.177

185/375

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ACATACTCAAAACTCTCATCCCTTCCTAAGTGTTTTACTACATTCTGTTATTTCTGTTTGTGAGGTCATTTGTTTATTG
CATCTGTATGATGGAAACACTGTATAAAATGACATGTTATTCCCAACTCCACATAGGCTGATATAACGTTGGTAGCTTA
AAATCAGCTTTAGTCCAGGTATTTATACAATGGAAATGGAAATCAGCATACGCCACAAATCAGGAGTTGATTTTTTTTT
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CTGCAACCTCCGCCTCCTGGATTCAAGCGATTCTCTGCCTCGGCCTCCTGAGTAGCTGGGATTACAGTTGCCCGCCACC
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GTAATCCACCTGCCTTGGCCTCCCAAAGTGCTGGGATTACAAGCATGAGCCACTGCGCTCGGCCCAGGAATTGATTTTT
TGTTTAATGGTGTTGACTGTCTACAAACTGATTAAGAAAATGTTATTAAAGTTGTACATCATGTCTGCAACCATTAAAT
TGTAAGTAGCACAAAAAATCTGGGGGAAACAGTCTTCTAGACTTTCCAGATGCAGCAAAGAAATTGCCTTGTTCCAACA
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TTTTTTTTGAGAAGGAGTTTAACTCTTGTCACCCAGGCTGGAGTGCAGTGGTGTGACCTCCGCTCACCATAACCTCTGC
CTCCCAGGTTCAAGCGATTCTCCTGCCTCAGCATCCTGAGTAGCTGGGATTACAGGCACCTGCCAGCACATGTGGCTAA
TTTTGCATTTTTAGTAGAGACAGGGTTTCACCACGTTAGTCAGGCTGGTCTCGAACTCCTGACCTCAGGTGATCCACCC
TCCTCAGCCTCCCAAAGTGCTGGGATTACAGGCATGAGCCACTGCGCTCAGCTCACCATCGCTTTTAGATAAGGAAACT
GAGCCCTAGAGAGTGGTTGGCTCGCCTCAGGCTCCAGGACAAATATGACTTAATCAAAACTATACTCCTGTTCTTTCAT
TCACATAAAACTACTTATCTAAGGATGCTGCAGCAACACTGCTGTCAGGCCAGAATTCAGTAAGTTTACAGCTGAGGCC
TTATCTATAGACCATTGATTTTGCTCAAGGAAAAAGTTACACAAACTAGCAATAGAGTCCTGACCAGGCATTACAAATT
CTACACTGATGTGTAAAAGAGGGACTAGGCACAAAGAATACATGCTTAGCACAATCTATCTTTATTATAGGAAAAGCAA
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TATAATGTGTAGGCATTGTCATATTAGAGAGACTCCTGGGAAATGCTTGGTCAACTAAAATTGTTAAAGAGCTAAAATT
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CAACTTCTGCCTCCTGGGTTCAAGCAATTCTCCTGCCTCAGCCTCCTGAGTAGCTAGGATTACAGGTGCCTGCCACCAT
GCCCAGCTAATTTTTTAATATTTTTAGTAGAGACGGGGTTTCGCCATGTTGGCCAGGCTGGTCTTGAACTCCTGACCTC
AGGTAATCGACCTGCCTTGGCCTCCCAAAGTGCTGGGATTACAGGCGTGAGCCACCGCGCCCGGCTCACTGACGCTCTT
CTTAATTTCTTCTGCCTAAGAAATAGAAGGTATTCTCGAGGGAAAAAAGGCATTGACAACTAGTACAGGGAAATTGATG
ACAACATAAGGCTGTTTGAGGGAAAAAACGTGGGATCTCAAAGACTGTTGGTGGGAGTGTAAATTGAATAATCTACTTTG
GAGAACAATTTGGCAGTTTCTATTAAAATTTAAAAATGCTGATATCTTATCAGTTAATCCTACTTCTAAGTATCTATTA
TTAAAAAAAATACTTGGCATGTATTCCCAATGCGTTTTGCAAGAATGTTCTTTGTAATTGTCAAAAGGTGGAAATCTGA
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CACAACTGGTAACAGAGTTACCTGGGGGAAGGTTGAGTTTGGGGATGGAGGGCTACAGAAACTTTAGAGTTCTGCAGAA
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TACAGCATGTGCCACCATGCCCAGCTAATTTTTGTATTTTAATAGAAACGGTTTTCATTGTGTTGGCCAGTTTGGTCTC
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GCCTGTTAACTATCTAAAAGCATAAAAAATAGTAAAACAAACGGCAAAATCATTTCCTCCTTCCTTTTCTACCTGTGTC
TAGTGTCTATTGAACACTTACTATATTCCCAGCATTTAAAGTACAAGAATCATGAGGCAACTGCTGTTTAGACTGAAGT
GTTTTCACCAGGGGTGTATCAAGAGGATTTCTTTGTCATAGGTTGATTTCCCTTATAACTGATTGATTAAATAATTACT
AGTTGGCTTATGCAATGTTTATATAATTACAGGATATAAATGGTGGTTATTAGCCTAACTAGAATATATGCCTTTATAA
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CAGTTATCCATAGATTGACTATAAGGCTAGGTATTGAGTTGGCGGATGCATACTTTCTTAATTCTTTAGATAATAGGTC
AATAGTGTGGTAGTGAATCTTGATGGTAAGTGTCCTTCGCATGTTTCATACTGCACAGTAGACTAGACTGAAAGTCCCA
GGAGACTGTGAGTGTAGCAAAGAATGAATCTTGTATCCCCAGATACACAAAGACCATGTCTTCACTGGACCCACAGAAC
CTAGCACATGGTGTGTCTTATTAAATGCTTGTAGAATAATTAAATAAATTACATTCAGAAAACAGTCTATAAGTCTTAA
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ATCTTTTAGAGTGTATATGCGCTTTTGAAAAAAATGGTTTATGATTTTGTTATGTAATACTGTTATATAAGTTATTTTA
GAAATTGCATGATTTATTATAATATCAGCAGTGAATTTAAACATTACACAAATGCCTGAATATAAAGAAATAGGATCAC
CTGTGTCCAAGTTAATTGCCTTCTTTTGCTGTATTCATTAGCCATCAATACCTACATAATCTAAGGCCAAGATACATAT
GGAGCAATCTATGGACAGGCAATAGCATCCCATGGGAGCTTTGGAAATTCAGGAGCTCGAGCCTCACCCCAGATGTACT
GAATCAGAATCTTTATGTTAGCATGGTCTCCAGGTGATTTGTAGGTCCATCAATGTTTGAAGAGCACCTGTCTGGCCTT
CAGTTGCCTTTATACATCTTCTTACCAACAGTACCATCTAGTCCAATATTTTTCTGTCCTGTTAGTTTGTCCATAGTCT
CCCAACAGGCTTCTGTCCACAAAGCTGTTATTTAGAGGCATGGCCCCATCCCTCTCATAGTCTTTGCTAAGGTTATGTC
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Fig. 6.180

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TTTTACATTTGGAGATACACCAATTGCAGAGGAGGGTGAAATTAGCCTGAGAGCAGGACCATAACAATAACCTGATTCT
AATGGAGGCTGTTACCAGGGCAGCCAGTACCCGAACTAAGTCATCATTCTGCTGGAAGCATTCCGTTGGCTGCAGTTC
AACAAAAAGAAAAAAGAACGCACTCCAGAAAATCAGGAAGGTTTGCATTATCTATCAGCTATGTTCAAGGAGATGTTTG
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TGTTTTAGAAAATTTGACTTATTATGTAACAGTGAATATACCTATCGTGTAGTTTCACTATTTTAGAAGTAAAACAAT
GCTTTGATTTCAAATATTAAATTAACTTAGGCATTGCTCTCCCACTCCATCTCCCAAGCTGTGACTCAGATATCAGAAC
AAAATTTTTTACCTAGTTAGGGAAAAAAGTTTCCCAAGTTTCAGAGCCATAAAAGCATCAGTAAAGCACCTATTTATTGT
AGCACTCTCTTGTTTTCCATCTTCAATTTCCTAAACATTGAATATCTTGTTTTTCATTTTTCTTCTAAAAGCATTTGTG
GTATATAATAACACCGCATGAAAGTGGAAGAGACTATATTGGCTCAGGATGCACTTGATATTTGTTTTCCTTAAATGAC
AAATGAAACTCCCAATTCAACACTAGGATAAGATTTTTTTTAAGTTATTAAACCTTAGTTAAGTTATATGACTTCATAC
TGGGAAAGTGAGTTCCAGAGAAGTTTAATGAACTTGTTGAAGGTGACTTTGCTAACTTATTACAAAATGGGGAATACAA
CTCTGGTTTCCCTAATGCCTTTTGTAAAGCATCCCAGATAAAAATCTAGGTTTCAGAAGATTCCTTATAGTTAGGTATG
AAAAAATGTTATTTCTGAAAATATGATTTCCTTATGAGAAACCATTGTTTAGATATGTTTAACTTTAGAAAAAAATTTCC
AACTATAATTAAAATCTATGAATTATAGACATGTTCAGTGAAATACACTGTCTCATAGAAACCATATTCAAAAAAACAGA
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TGAGTTTTGAATAGATCACTATTGTGTTATATTCCTCTGAATTTAGGATACCTGTTGGATTTTGTTCCTTATCATGTCT
GTTATGTGGACAAAGACAACAAACGTTTGTTTCTTCTTGTCAATTTCATTATGCACAAGGATTCTGAAGGTCCACCCAA
GATCTTTCGTGCACTAAATGTCTGGTGTTTGTAATACAATGAACAATTTTGGGGAGTTTGAAAATCTTTTGATTATGAC
CTTGAGGATTAGAGATCATTTGGTGTGGATGACTTCGAGGAAGTTATTTAATTTTTTTTTTGTATGTCAGTTTTCTCAG
ATTTAAAATGTGGGTAATAGTAACCACCTTGTTAGAATGATTGTAATGATCAAAACATGTAATAATGCAAAGTGCTTAT
GCTTAGTGCACTAGTGTTTTGCACTTACATCTAGTTATTTTTTTCTCAGTAATTCAATAGGCTAAGGGATGATCTAGTA
CAGATTGAGATATGGGATTTTTGTTGTTGTTGTTGTTGACAAGTTGCAGACAAAATGTTTGGTTATTCCTCTCACTGAA
ATAAAACCCAGAAATATAGGGTATTATAATATGTTAAACATTTAGTGGCTATCAAAACTTATTTTCTTCTTTGAAGTCA
GAGTAATATTTAGAGAGGGCATTCTGGGGTTTCCTTTTAGCAAAATAATTAGAAGTAATTTCCCTTAGAATTTTTAGAA
TGACTATATTAGGAGGAAAGGGAAGGTTCTTACCTAAATTGTATTGCAACTTTTCTCTGAAATAAAATAAACATGCTGT
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CACTTGAGGTCAGGAGTTTGAGACAAGTCTGACCAATGTGGTGAAACCCTGTCTCTACTAAAAACACGAAAATTAGCCC
GGCATGGTGGTTGGTGCCTGTAATCCCAGCTACTCACTACTCAGGAGGCTGAGGCAGGAGAATCACTTGAACCCAGAAG
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TGTTGGGAGTTTTATTTATATTCCCTTCTAAAGTAGATATTTTACTTTTTTGGAAATCTCTGCCTACTAGCCCAGTACC
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GGTAAACTATATGACCTCTGGCAAGTTATTTAACCTCCTTATGCCCTCATCTCTAAAATTGTAGTACGAAAGCAACACC
CTAAGTATGATTGCAAGGATTAAGGGAAATAAAATGCAAAGTTCCTGGTACATAATTGATATCTACTAAGTGTGAATTT
CTCTAAGTTCCTCATATCACATAATGGTAGATTTTTATTTACTGGGTGGTTATTCTGTACCAGGCACTCTCCTAAACAC
TTATATCATGGGGATTAACACATTTAATCCTTGTAGCATTCTTTTGCAATAGCTACTATCATCATCATCCTCATTATTA
TCGTCCTATTTCACAGATCTGGAACTGAGCCACTGAAAAGTAAAGTAACACTAACTAGTAACTAGTACTTCTGGGGTTC
AAAATAAACTTTTCATTTCTAAAATTAACCCATACCTTCTTATCCAGTAGACAACATAACCTCTCCCACTATCAGGTCA
TTTCCATGGCCTATATCCTGTGGACAATTTGAAAGCGATTTCCTTTTAACCCTTGTTTATCTAATTAGAACATAGTTTG
GAGAAACATACAATGTTTTCCATAGCTGAGTATAAATCATTTTCTTCCATCCTTTTGAATTAGCCACATCATTGCTCTC
TTTCCATTGTGGGAAATCTAGCCTTGCTTGTTTGTACCTAGACAAAAGCACACTTTTTGTGTCAGAACCTGTGAAAATT
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Fig. 6 : [8]

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TTATCTTCCTTGTTACCACCTATAAATGATTAATGAGACAAAACATTTTCAAGGATTCCAAATAATCTAAAAAATCCCA
CAAATTTTGAAAATCCACAAAGAGGTATCTAGGAAAACTTTCATGAAGTTAAGAGATGTTA1. .GCTGTTCTAGGCGTG
GGTGACTTTATCAAAACATTTCGCCCACTTTCAAATTTCCACTTGTTAAAACCTGAGGCCTTATTTGGCTGTCACTTAC
TGTACAGTGTTTTTCAGACAAGTTCCTGGCAGCTAATTTTAAAGAGATTTTGTGAAAAACACAAGCTGTGGAGTGCTCC
TGTTGGAAGAGCACCCTCTGACCCTCTGCCTTAACATAAGAAAAAATTGCCAGTCTGTCCAAACCTCACCCACCTTTGG
TTATCTTCAGAATCTTACAATTTGTGTTTCCTAAGGCAGCAATTTTCAAGCATGACTTTGAAACACTGCAGGCTTTTTC
AAGTTATTTCAAGGACTGGCCAAGAATTCTCAAAACAAAATTAATTTTGTATTTTTAAAAAGCAGTATGCATATCATGT
ATCATTTGGCAGATTTAGGAGGTTAACAGTAAGCATTAGGAAATTGAATCAATAATAATAATGGGATATGGGGACCTGT
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TCCTTTAACATACACCAAAGCAATGGGAGATCTGCAGAATGCTTTTGTCCATGTTTAACCATAATGATCAATGCATTTT
GAATAGAATATTGTTATATCTACTATTTAGCTTAATTGTGATCATTAACAAAGTGCCCAGACTCTCTGAACATTCTACA
TGGCATACTATTTGTGAGAAGATCAGCATGTAAATAGTTTACTCTTTGTAAGCTCTTCTACAAGGTGGCTAAAAGCAAG
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GGAGTCTCGCTCTGTCGCCCATATCATTTTTAGTATAATCTCTTCCTTTACTTGGTCACTAAGAATCACTTTTCTCTAC
TTAGAGCCAGACATTTAAATATCTCCAGCCTCCAAGATCTCGTCTGGGTACTTGGGCTGCACTTTTTCATGTCAATCAT
TTTGCACTTGTCATCTATCCTATTAGGCTATAAGCTCCCCAAGGGCAGATATAGAGACATAGTATCAAGCTAACCTTTT
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GATCATGGCTCAATGCTGCCTCAACCTCCCAGGCTCAAGCAATCCTCCCATCTCAGCCTCCTGAGTAGCCTGGGACTGC
AGACGCATGCCACCACACTCAGCTATTTTTTAATGTTTTATAGAGACGGGGTCTTGCTATGTTACCCAGGCTAGTCTTG
AATTCCTGAGCTCAAGTGATCTTCCTGCATTGGCCTCCCAAAGTGATGAGATTACAGGTGTGAGCCACCACACCTGGCC
AAGCTAATCTTTAAATATCCTAGAGCACCCATTATAGTGTCTGGCAAAAAATAAATGCCCATAAATGTTTGTTGAATGG
AACTGAGTCTTCAGAAACTAGAGATTTTTTCTAAAATATTTTTTTTTGCCATTTCAGTTTAATACATATTAGTTTTTCT
CACTTATCCTATCATCTGAATTTTTTATTTCTTTGAACCTGAACAGGCTTTGTGTCTACAGACTGCTGAGCCTTTAATT
TTGTTGTCGTTTGTTTTGTTTTGTTTTGCACAAAACTAAAAACATGTACAAATACATTTTTAATTGCATTTGTTCATTC
TAGTTTTTACTCGTAAAGTGCACTTCTTTCTTTTCTTTCTTTCCATTAATTTCTGAACCTTCCCTTCTTAAACTTTACC
CAGAGTTTTCTCATCATATACGCATTTCATTTTCATTGTACTTTTCTGCATCTGACTATTTCATTTTCCAGTTGTTCTG
GTATGTTTTTTCTAAGTATTGTCCAATCCTTTCCATTAAGTTACAAGCTCCTTGAGGGCAAGCATTGCACACCTGTTTTG
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TGTGGGCATGCTAACAGGAAAGATGGACCAAGTGGTTGACCCTGTCCTTCTGTGAATAGAACCTCAGAGTACACTTTTC
TGTAAACATTTAAAAGACAAGGAATTGGAACTTAAAATGGAGCAACCAGGGGGGCAGGGAAACAGAATAAACTTTTAAA
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TCCCTAAGTTATTGGGTTTTTAAGAAACATAAAATATTCCTAGATCGTTAGAGATGTGATTTGATTTCAATCCCACATG
GTCTGTAGCATCACTAAGCTACTGGGAACTTCCCTATGCAAAGGTTTGAGTTTCTTTCCAATGCTGTTTAATGTCTGCA
CTTCTAGTGTCCACATAAATGAGATTTAACAGGCTTTACTTCATACTTTCTTGTTACAACATTTTTACTTTTTTTCCAA
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AGGCAATAAAGGAAATCACGAAAAGCTAGATGTATTCCACCCTCTTATAGTTGAATACATAAAATTGTATATATCACAA
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.ACAGTACCTGAAGTGGATAATCCGCATTGTCCAAACCCGTGGCTGAACGAAGACCTTGTGAAATCCTTGCGAGAAAACC
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CCCCAGGCTTCTGCGCAGAATGCTTCTCAGCAGCAACATCCCCAAACAGCGGCGTTTCACGGTGGCACATACATGGTAA
GGTTTGCGCAGACCTCGCAGAAATGGTGGCCGTTTCCCTTCAGGGAGTCCTGCTCTCGCGCCTTTAATTTTGGGGGTAG
GGAGGGAGGACCATCATCATTCCTCACCGGGGATTTATGGCCAAGTGATCTCCCTTAACCATAAATCCCACAAATGTCC
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GTTCCTACAAGAAAATTGGATGCCAGATGTCAAATAAGAATGCAGTTTTATTCCCATACGTACGTAGTGGCGTTTCTTT
GATTCCTTAAACCAATTTCCCTGCCAGCTTCCCCAACAGCTGAGGAAAGAAAGGACCAGAAATGAGGGGAGGAAAATCA
```

Fig. 6.182

188/375

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AAAGCCCTTCCTCACTCCTGACTTTGATATACAGTATTCATAATCTTAAACATGCCTTTCCAAAGGAAAGAAGAAAGGG
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Fig. 6. 183

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TAAAATGTTACAGATAATGTCTAAAGCTGTCACCCATCCTATTCTTCTTCCTTCGCAGAGGATCCTGAAATCAGTGTGT
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ATCAGTCTCTTCTACTGTAAATAAGCAAAACAGAAAAATCATTTATGTGCAGTTTTAAGGACATAAATACTCTCCAAG
TATTTCAGATGAGGCACTTATTCTCAAAAGGAGATCTTGAAAAGTTGATCTGAGAAGAAATTACATGATTTCATTTTGG
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Fig. 6.187

AATAAAAACACTTAGGTTCATAATAATATAGCTTTATAACCTTAGATTAATTATAACATAGACGTCAATGATTTAGCTC
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TCACTTACTGATATGCACAGTATTAGTCATGGCAAAATATTATTAGCATTTGAAATTGATGCTCATCCCTCTTTGTTGA
TACTGTCTCATTTACTGGATTAACTTTATTCATATGAGATCTCCCTTTTCTTTTCATTGTCTCTGTTTCCAGCACCTTA

Fig. 6. 185

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ATTGAAAGTACCCCCGGTTACATAATTATTCCTTTGACCTTATTCAGTATGTTCCCTGCACCACAAGTGTCTCTCCTCT
GTTCAATTCCAACTGTCAGCTCTTTTAATAAACCCTACTCTTAAACCTCATTTCCCAGGCTCAAGTTCCATATGCCTCC
TCTGGACATTTCCACGTGGAAGATCCACTCTTAATTTATATTTAACACGCACAAAACCAAGCTTACTGTGCCCCCCTTT
TTGATTTTTCTACTTCTGTTTATCTGGCCATTATCTTCCCAAAAAACCAGAAAAAATCATATTTGACTTCTGCCTTGCC
CTCCTTCTGCCTAGGCCTATGGATTATTTCACCTATTATCTCTTGCTGTTCTGTTCCCTCCCAATACCACCAGGTATTT
TAATACTTAGATTTGTGTTTGATCTATTTCAATAACATTCTAACTTGTCTTTGTGCCACTAGCTCCTTCCCACTAAGGA
TGTCCTACTTATACTGTCTCATTAATTACCCTACAGTTCTGCTTTGATTGTCCACTATCTATAGCAGAGGTGAACAAAA
GGACCAGAAAATAAATATTTTAGGCTTTGCGGACCCTATGGTCTCTGGGCAACTATTCAACCCTACCACGGTAGTGCA
AAAGAAGCCCAGATATACAAATGGACATTAGCATGTTCCAATAAAACTGTATTTGCAAAAACAAATAGGCTGAATTTCG
CCCACTGGCCGTAATTTGCCGATCCCTGCCCTGTAGGATAAAGTTCACACCTTTAACATGGCATCTAAGCATCTGTCAT
CAGTTCTGCACCTATCTCTGCAACCCTATGTGTTATTGTGTCCTATGGATACTTCACATCCTTCGTTTTGAAGCCGAAT
CTAGAGCAGGGTTGCTCAACCTCAGCAGAACTGATGACTGGGCCAGAATGATGCTTTGTGGTGGGGGCTCTCCTGTGCA
TTGTAGGAGGTTAAGAACATCCCTGGCCACTGCCCACTAGATGACATTAATTTCCACCATGAGGTGAAAATTCAAAATG
TCTCCAAGCACTGCCAGTTTTCCCTTGTAGGGGACGAGGGATTGTTTTCAGTTAAGAACTACTGCCCTAGGGAGTAAAT
TTTGTGAAGGGGAGAACCAGTTCTACCTCATTCACAGGTGTATCCCTAGCACCTAGCACAGTCTTGTGATATACAAAGA
GGTTCAGGAATCTATTGCATAGTCCAACTCTGCTACTCACTGTGGGATCCTAGGGTTATTTAAATTCCACAAATCACAA
TTAAGTTCATCTGTAAGTGGAACTATTACTAATTCATACATGTTTCATAAAATTATTTAGAAGTTCAAATGATATATTT
AACAAAAAGACCTCTTGAAAGACAGTAGTACTTTCAAATAAAAGGTAAACATACAATTGATAGAACACTGAAGGTGATC
TGAGCCTATTGAAATCTAGAAGCTTAATTTGACTCACATACTCTGTCTACACTGTACAAATTACTATATGGGATGTTAT
TGGGGCATTGTCTTTGTTCTCAAGGATTATGACCTAACAGGTGTTAGAATATGATTACTCAACCATAAAACAATAGTTG
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CTGCAACTTGATTAGGTTTAGGTTTTATTATCACATTTCACAAAGAGAGAGGGGCCTAGGAAGGAATTTGATATGAATG
TGAAGCATGGGAGATTTCAGCAGGGAGAGAACTGCAGTGAAGTGTGGAAGGAGGGGAAGTGACAGGAGGTGTTTTCTCA
CCAAGGGAATAGTGTGAACAAATATGGGGATTACATGGGGTTTAGGTGAGTTTGAGTGCCTCTAGGGAAGGTAGGTTGG
AATCAGATTTTGAAGGATTGCTAACATCTCTCTCTCTCTTGCTCTCGCTTGCTCGCTCGCTCTCACATCACACAC
ACACACACACACACACACACACACACACACACACCAGGGATTTATATGTATAAAAGAAACATGATGATGTTATCTTTCC
TAACAGCTTCACTCCACATAACAGGAAGTGACATTTTGAGTTCTTACCCCATGCTAAGCTTGCTAAGCCACAAGTATGT
GCACTGAACACTTTTCAATAGTTGGACTTTGAGACTCAGTTCAATTAGACCAATCTATACCATCTTTCAGCTAGCTACTG
CTCTTCTACTTTTCCCAAGATAATCCTTCTGAAATAGATACACAATGCTGTTTTCATTCAAATCAGATGTGTTAATTCT
ACTAGTGGCACTTGTGTTCTGATGCATAGAACAGAGGCATTATTTACGTGAGGACTGTGCCTTCCTTCTTACTGAATTG
CAAAAGTATACAATTCTTAGAACACATCTAGCAGTTCTGAGGGGCTGGACACAGAAAACCATCTTACTCTGAAGGTAAT
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TAGTTTGAGTGTTTCCTGTGTACCTATTGTTCTTCTGGAGATAGAGAACTGAATGAAATAGACAGTCTTCCATACATTG
GGGCATACTTTTTTTTTTTTTTTTTTTTTGAGACAGGGTCTTACTCTTTTGTCCAGGCCGCCTAGAGTAAAGTGGCA
TGATCACTGCTCACCTAAGCCTTGACCTCTTGGGCTCTGGGTTCAAGAGGCCTCCCACCTCAACCTCTCAGGTAGCTGG
GACTACAGGCACATGCTACCAGGCCTAGATAACTTTTTATTTTTTGTAGGGATGGGGGTCTCACTATGTTGCCCAGGCT
GGTCTCAAACTTCTGGGCTCAAGCCATTCTCCAGCCTCGGCCTTCCAAAGTGCTGGGATTATAGGCATGAGCCACTGCA
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CCTGGCCAACATGGCAAAACCTCATCTCAAATAAAATTACAAAAATCAGCCAGGCGTGGTGGCAGGCACCTGTGATCCC
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CTGCACTCCAGCCTGGGTGACAGAGCAAGACTCTGTCTCAAAAACAAAGAAACAACAACAACAGAAAAGTGATAAACTG
TGGCAGATGCTGCTCATAGGTCAACTAAAATAAGAACTGAGAATTGATCATTGCATTGGCAACATGCAGCCAATTGGTG
GCATACAACACAAGTCGATCAGTAGAGTAGTGATGATACAGGACTGACTAGAGTGGAATATAACAAATTTCCCCAATGC
TTAAACCAACAAGCATTTATTATTGCATACAGTTTCTGAGAAATCAGGGAGCAGCTTAGCTGGCTGGTTCTGACTAAAG
TTCCTAGAAGAAATTGCACTCAAGTTGTCAGTGAAGGCTATAGTCATCTGAAGCTGTATCTGGGACTGGAAGATCAGTT
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Fig. 6.1186

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TTCTAAGATGGTGCTGTCCGCAGGCAGCCTTGCTTTACACTGGCTGTTGGCAGAAGACTTCAGTTCCTCACTGTTTGTG
CTTCTCCAAAGGGCTGGGTGTTCTTTTTTTTGTTGTTGTTTTAATACTTTAAGTTCTAGGGAACATGTGCACAGCGTGC
AGGTTTGTTACATATGTATACATGTGCCATGTTGGTGTGCTGCACCCATTATTAACTCGTCATTTACATTAGGTATATC
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TTCTCATTGTTCAATTCCCACCTACAAGTGAGAACATGTGATATTTGGTTTTTTGTCCTTGCGATAGTTTGCTGAGAAT
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TATATGTGCCACATTTTCTTAATCCAGTCTATCATTGTTGGACATTTGGGTTGGTTCCAAGTCTTTGCTATTGTGAATA
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GATGGCTGGGTGTTCCTACCTCAGGCACAGGAGTCAGTGATTCAAGAGAGTGCAAAGTAGAAGCCACAACGACTTTTAT
GACCCAGACTTGGCAGCAACACACTATTACTCTGCCATATTCTTCTGCTCACATGTGCCAACCCTGGTACAGTGTGAGA
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GATCATGAGAAACTGGGGACAATTGCAGGAGCAATATTCTTGAGGAGGCTTTGGGGGAATAGGGGTAAGGAGAAGTTCA
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GGACAGTCTGAGAGTATAGGAGGTCAGTATCTTGATAGATCTGGTTTCAGAGTGTATGAAAAGTGTCTCTTAATTCCTT
CTATTTTGTGATAAAATAATAAGCTCATCAATGTAAAGACAGAGAACAGATGTTGGAGGTTTGAGGAGAGAGAAGGTGA
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GGATTAGGAGGGGAGCAGATTAATGATGTGAGTTACAAAGACATTTTAGGGCAAGGAAGAGATCATGGTCTGGAAGCAG
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GAGGGCTGAAGTAAATCCGTATTCTCGGTGGGCAGCCAGTAGTGTCAGAGCAAGATGGTGCAGGAAAGTCAGAGAACAT
GCTGAAGATACAGGAGCATAAGGAGATTTTGCTGATGACAGACTGAACGCTAGGAGTCACGCTGGCGAGTTTGCAGGAA
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TGAAATGTATAGCTTTCTCTTTGGAGTTTGAATTGTTGGTTCTTACTGCCTGCTTGCATATTTATATTAGTTGTACTGG
CTGGCAGTTATTAGGAAATACGATCTTCTGTGTCTTAGGAGTGCGCCCTGCAAGCAACATGTGCAAACATTTTTTTCTT
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GTCCAGAATTGGCTAGAACATGCCTGTCACTCCCAGCTGACTCTGATGACGCTCTTGCCAACGTAGATTTACATCAACA
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AGGGCCACCTGGAAAGTTAAGGTCATCAGCTTTTTGTTATAGTCATTGGATTACTGTGATGCTCAGCAATAACAAAGTT
TGACCTTGTAAAGTGATCAAAAGTTGTGGGTTACAATATAGGAATATTAAAGTATATTCTGCCATGATGTGTCTTCTGT
TGTAGTTTGTCTAGAATTAATATTACAGTCACAATTAACTCCAAAGTGTTCTATACTGCTTGCATACAGTACAGTGGGA
AAGAAGAGATGTCAAAATCAGAGAAATAAGATTCCTCCATTATGAATTAACATACAAAATCTACACTTAATGCATCAGA
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Fig. 6.187

ACTAAGATGCAATGTTTTAAAGAGATTGAGTTGACAGGTCAATCAGATGGTAGGACTGGTAAGAATTTGAGCAGGCATA
AACAATTGGCAGTATGTATTCAAACACTGTTTATGAGAGTGAGGAACTATTGGTTAACAAGCTGACATTAAGCATTTAG
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GTTAGCAGTGTGCTTGTAGCGTATTACACTGCTGTATCGCTGTATCACTGCACCATCTCAAATATTGAGTTTTTTTATA
GTTAAGTATTTATAAAATTAATATGTTGGATAAAATCTAATTTATCTCAATTGATTAACATTGAGGCTGCAAATATTTT
TATAGTGTAAAAGAATTTTCTATTGTGAATTGTACATTAGTGTCTTATTGAGACATTAACATTTCCTGTCTTTGTTGTT

Fig. 6.188

GTTGCTGTTGTTGTTGTTGTTTTGACACAGAGTCTCATTCTGTCACCCAGGGTGAAGTGCAGTGGCACAATCTTGGCTC
ATTGCAACCTCTGCCTTTTGGGTTCAAGCGATCCTCTCATCTCAGCCTCCCGAATAGCTGGGACTACAGGTGTGCACCA
CCACACCCAGCTAATTTTTGTATTTTTTAGTAAAAATGGGGTTTCACCATATTGCGTAGGTCAGTCTCGAACTCCTCACT
TCAAGAGATCTACCTGTCTTGGCCTCCCAAAGTGCTGGGATTACAGGTGTGAGCCACTGCACCCAGCCCTTTCTCCCTC
ACACTTTTGGAAAAAATATTTTTTCTCAGTTTGTTGCTTGCCATCTAATTTATTACATTGCTTTTTAATATACAACTATA
GACAAAAGAATCACTTTTCTTTTGGAAAATTTGATCTATGACTTTCATTCATATAAAAGCTTTCCTCACCCTCAGGTCA
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TGTTTAGAATGAACATGAATGATAAACTAATAGGTATTATTTTACAAATTATAGAATATTAAAGATAAATATACTCTAG
AGTATAGAAATATATTTTATATTATTATGTTAAAATATTAAATATTTCAGGCATGTAAAAATAGATGATAGTGTAGCCA
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TCACAATACCGTGGCCTATTAGATTGCTAAGAAATTATGTATGTCAACTCTTTCTTTATACAAATATTGAAAATGATAG
TAAGATTAAGGAACTTGCCCAGGTCACAATTTTTTTCTTTCAAATTTCATGGCTGATTTCTAGGCAAAAATTTCAGGGG
CCTGTCCAATACGCATGAATTACCCTACTATAAACAAAGCTAGGGAATTTAATACATACCGAAAGGCATTTTGTAATAT
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CACTTCCAGGAATGCGACTTTAAAAGCCTAGGCTTTATGTTATGTCATCAAAGTCATTGTACTCAATTAGACAATAGTG
CTATAATTAATTAAATGTTATTATGTAAATGCAGAATTACCAAAGAGAATTTCAAAGAGTATAACAGTATTAAGTCGTT
TCCTTAGTAATTTCTCATTGCTGTGAAAAACATCTGTAACCTAATACATTTATATTCATTTTTCATATTATATTATGCA
CATATTATTTTTAAATATTTGTTAACAGGTAATTTTCTGATTACCAAAGTAATACTTTTGTATTAAATGTTTGAAGACT
GCCTCCAAAAATTGCAAAGAAGAAAATTTTAAAACCGTTATAATCCTACCTTTGAAATAATTATTCTTCAAATGTTAGA
TATTTGGAATGTTTCCAGGTGTTTACTGGTATAGTTAATGCTGCTGTGAACATCTTTATTGCACATACCTTTTAACCTC
TGTGGAACTGATTGCTTCTTTCATGATTTCTCCTATATTTTGTCAATCATTTCCTTTTTTTCTTTTTCTTTTTTTTTTG
AGACAAAATCTCACTTTATCACCCAGGCTGGAGTGCAGTGGCATGATCTCGGCTCACTGCAACCTCTGCCTCCCGGGTT
CAAGCAATTCTCGTGCCTCAGCCTTCCGAGTACCTAGGATTACAGGTGTGCACCACCATGCCCGGCTAGTTTTTGAATT
TTTAGTAGAGACAGTGTTTCGCCATGTTGGTCAGGCTGGTCTCAAACTCCTGACCTCAGGTGATCCACCCGCCTCAGCC
TCCCAAAGTGCTGGGATTATAAGAGTGTGCCACTGCACCCGGCCAATTGTAAGAATTATTTTCAAAGGAATTTATATCA
AGTTACAGTGCCCCAGAATATTCTGTTATTTTAGCTGTATTGAATATCATAATTTTCTTAACATGTTTTGTCTTTAGAT
GGTATAAAATACTATCTCAATATTATTTTCCATTTGTTTGTTTACCCTTTATATTTCCATTTGCTTGAACTCTCTGCTG
GAATCTTTCAATATCTACCTGATGAATCAATGTTTATCTGTTGGTTTGTTTTCTTTTTTTTTAATTAATTAATTTATTA
TTATTATACTTTAAGTTTTAGGGTACATGTGCACAATGTTATACATGTATTTTTATTTCTTTTTGTCACATAAATTTGAC
TCTTTATTTTAAGCAACTATTTCTAGTCTTTTGTTTCTCTTGTTTTATTAAATTTTTATTGTAAAGAAGTTTTGTTTTG
GCCAGGCATGGGGTGGCTCACACCTGTAATCCTAGAACTTTGGGAGGCCGAGGCAGATGGATCATTTGAGACGAGGAGT
TCAAGATCAGCCTGACCAACATGGTGAAATCCCGTCTGTACTAAAAATATATTAAAAATTATCTGGGTGTGGTGGTACAT
GCCTGTAATCTCAGTTACTCAGGAGGCTGAGGCAGGAGAATCACTTGAACCAGGGAGGTGGAGGTTTCAGTGAGCTGAG
ATCGGGCCACTGCACTCCAGCCTGGGCCATGAAGCGAGATTCTGTCTCGAAAAAAAAAAAGTTTTGTTTCTGTGATAGAT
ATTTTGCTTAGCATGCTTGCTTTTTATATTTAGAAAAACTAAATCTATTCCATCTAAAATATATTTTGGTGAATAACGC
AAGATATGTAATCTCAGTGTTATTTATAAAATGTACCATTCTCACTTTCTTATTTTGTAAAGCTTGTTTAATTTCAGAG
TTTGATATATTGAAGAGTCCTATTCCACTGACTTAAAAAATTGAGGCAGAAGGATCCCCCTCAAGTGTCACATCTTAGA
ATTTGTTTGATGGTACTTCAAGGCAGTTGGTGAATAATTTAGAACTCAAACTTTGGGCTGCAGATTGCCTGAATACAAT
TAATAGAAAACAAATATTTCCTCAAATTACATTCTTTGACATTAGTAATCATTCCTTTATATACATCTCAAGTCTAAA
CTCCCAATCTGTTTATATGCAGAGATTCACAGCTTTAAGATTTATGTTTCATAACTGCAATATCACTCTATGATACATT
AATGGGATTCTGTACTCAACTATTCCATTGGCATTCAAGTGAATAATTTTTATACAAAACTTCTTCAGGAGACAGGCCC
AACTGAAGTGTATCACTTTAAAACAAATATCCTATGGGCAATAGATAAATCTGATATTTTTCTGAGTAGAAGAAACATA
AAACCTCAATATAGGATTAATAGGGTTCAAGGGGTTTTATAAGCACAGTGCTTGTGAAAGTATGTAATTCCTATTAAGG
CTTGCATTCATGAGCACATCATGGTATATGCTCTCTCTGGGAATATGTAAAGCCAGTTTAAAATTCAATTACAGACATT
GCAGTCTAGACCTCCGCAAGCCTTTGCACGTGACTGGACTCACTCAGGTTAATCTCTGAACCAGTTGTAGAGAGCTTTT
TTCATCTCTCCCTCACACATTTCCAAACTCTTTTCTTTCTGTGTTTTTTAACTGTGACTTTTCTTTACAAAAGGAAAAG

Fig. 6 189

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AGAATTTTTTTTAAGTCCATGACATATCCAATGAAATCGAAATGATTTAATACATGGAGTTATCTTAATATCTTTTGCT
TCTTGACAGTTCTCTCATATCTATCTAGAAAATATCTGGCCAAAAAAACCCACTTTACTTTGTTTATGAGATATTAGAT
TATTTTGTACATTTTCACATTCCAGGGCTACAAGGAGACCACAGTAGACAAAATCAAACCGTCTTCCTTTTCTTTATAG
GCTTGTCTCAGCAATGCAGCACTGTAATGTCTCTTACTTGAAGGAACTTCATTCTATTGTAAATTCTTAGGACAAATAG
AATAAGGAGATAGAAGAGTTGTGTGAGTTATAACTTATATGTAATTTTCTGTATATATTTGGAAAGTTCAACATCAGAT
ATTGAGTCTATTTCAGTCTGTTGCTGTGAATAGAATACACTATTTTATCCTATTGGTTCTTTTCTTAAACTGTCAACCA
TGGTTTGAGCATAGTTTAAGAAAAAAAAATGAATGCTTTAAATAAAACCAGTACCTGGGTTGGAGGTGGCAAATATAT
ACAAACCTGTGTATGTGTGTTTGGGTGTGCGTGTAATCTGAGGATTATAGCCCTGTATCTGTTTCAGGGAGAATTTTTT
TTACATCAGACCATGAAATTGAAAGGTGATTCTGATAATCTATTTCATTAAATAGAAGTTTATGATACCATAGACTCTG
GAATAAAAAATTCCTTAAAATTCCCTATTTAGATTAATGCAGATTTAATCCTTAACCCATTGGTTCAAAACTCAAGTTT
ACTCTTTCAGTAATAACAAGAAGTGGTTGTCAATAATACTCTCATTAAAAATAATTATTTCAGCATTTAAAAAGTAATA
AAATTGGTATTTTCTAACTTATATGCTTAATACTCATCCACAAAGGTTAAATAATTAAGAAATTAAAGACTGTGAAGAA
GAAGGGAAGAAAGAGGAAGGAAAAGATGGTAATATGGATGCTATTACCATTCACAAAAACAACATTGATGATTAGGCAT
ATGTTGACATTTCAATTTTTTGAATCTTGGTAAAATAGTCATTTTAATAACATACTTACCAGATTACTTAAAGTAACAT
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TAACACTTTCCAAGCTTCCTTCTCTTTTGTTAGATTATAATTCATGTACATCCTTATTGGGGTATGGGAGACACTGGAA
CTTATTATACTACATCTCATCCATTTGTTCTGTCTCCATGACTGACTTCTCAAAATAATCTAAACTCTTCAGTGAGCCC
TTTAAAGTCATGGTTTTATCTATATCTGTCCAGGATTTTATCTCACGTTTTGCATGTATCCCAGCTCAAAAGGATTAC
TTTTCTGAACTTTCATTGCCCTTGGATTTTGACCTTCCATATACTTGAAATTCTCTCTTTTTAATATCACCACCAATAA
AAATCCATCCCTCAAGACCTCACCCAATTATTATTTCAAACTCTAAATGCTGATCAACTATGTGGAGCTGACCTTTTCA
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TTTGAATTCTCCATTGTGCAATTTCTAGCTCACTGTCATACAGAACATTACTATCATATTTAGAGGCTGAATAAATCAT
TAAATGGATACCATCACTGGGCCCATGGAATTGACTTACTTAACTCAAATCCATGTGAAAGTCACTTAATAAATATTAA
ATACCATCAATTCTCAATTATTCATGGGGTTATTATTCACAACTTACTTGTGATATGTCCTTAAAGGTCATATTTACAT
AATGAATGATTCTGCCTATTTCATTGGCATAAATAGGTTCATTTAAACAGATTTCATTATTTAATTAGCTCCAATTTTT
CCTTTACCATCTTTTTTCAGAAGGCAAGATTTTGCTTTTAGCTGTCTTGCCTAAATCTGAAAGCTCATACTGATCATTTC
CATTTAATAAGGGAAGATGAAAAGGGCCGGTCAGGGGACTGGGATCCAAAAGGTCAAGAGTTTTCCATTCCTTCTTCCA
TCAGATCTTCATAGTGGAGTTTTGTTAATTTGTGAATTTAGTTCAATGAGCCAAAAAAAAAACTTGCTGATCAAAACAAA
GCCCCTTTGGTAGGGAGGAAATTCTGTCTCTTTGTTTTCTCTTCCTTTCTCTCTGTGCGTGAGTGAGTGTGTGTGTGTG
TGTGTGTGTGTGTGTGTGTGAATGTCTCCTGACATTTGAAATTCACAGTGCAATTTAACAATAAAATAATAAAATATAG
CTTTTTACAAAAGGTGTAAGCATAGATTAATGTACAGGTTTTCAGAGAAGGACAGGTTAGCTTGCTTCATTTTTAATCC
TTAGGTAAAACAGAGGAAGAGGAAAGTGAATTTTATTGAAATTCACACTATGTACAATGAATTACAAGAGAACTATAGG
AAATGGTTAGATATGATTAAATAATAATTATTTTGAAATTAAGATAAATGTCAAATGGGAATTATTTGCTTTTTCATTA
GGAACTGCAATGATGTTTGCGCAGCAGTGATCTATCTAGACAGATTTTGGAAGATGTCAGCAATTTTGTTTAGTCTGTC
AGCATTTGCAGTGGCAATGATTGGGATTTACTTGATGCATTCACTCTGGAAAAAACCTTTTGTGCACATTAGCATGACA
ACTGGTGTCCATGAAATACAAGAACAACATTCAGTTATAATGCACTTTACACCAAATTAATGAATAGTTTAGGAAGGGA
AGCAAGAGCTGACTTAGATGACTTACAAATATCTGCTTTTTTGCATTTTCTAAGGTAGATAATTTGTGTATATTTACTT
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AATAAACTTAGGAAGAAGGCGGTTGCTTGAAATATTGTAGTCTGGCACCAGCTCCTTCAAAATGGCCATGTTTTAAAAG
TATGGCAGGTGTGCTAAAAAGTCTTTATTGCAAAATCATTAGAAAGTCTTTTTTTTGTCAGTTTAAAATGCATTCAGCTA
TAAGCAATAGAAAACTAACAATGGTTGAATCATCTAAGGGTTTGTTGGTCTTGTGGAAAAGGATATCAGGAGGTAGCAA
ACTAGGGCTAGTACAGTGATTCTATGATACCCTCAAATATGTGATTTCCTTTTATTCTTTCTTCTGTGCCAGCCATCCT
CAGCATATGGGCCTGTCCTGTGCTTGTCATCTCACGACCATTGGCGGATACTGCTTCCCACGGCACWATCTCCTCATTC
CAGGGTAAAAGAAGAGAAATGGAAAGGAAGTAGGAAGGGATGGTGTCTGTATCAGGGAAACAAGGATTTCTTAGGAATC
CTCAGCTTATATAAACCCAGCTACCTTGCCCCTCTCTGACAACCTGTTCTTTGGCAAGAACAATTTCATATACTTGCCC
CTACCTTCCAGGGAAGCTGAGGAGCTTGTTTTTTTTAACTAGACATATTATTGGGTGATTAACATTAGGCTTATATAAA
TAAGAATGAAAGGGAGAATGGATATCGTATATACAATCAGCAGTGTCAGCCACAGTCTTATAGTTAAGTGGACACCACT
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CAGATGATATTACCTGTATACTATTATGTTAGGGGTATAATACCATAAATATGCTAAATTCAGATTAATAAAAAGTTTT
TCACTTTCTGCACAATCATGACCTCCTTTTATTAAAATAGACAAATAGTCTGGGGTTAGTCACACAAAGTCTATATGGC
ACATGACTACAATAATTGATCTGCAACCTCTCTGTTCTTTAAAGTGATTCTGTGATATCAAGGAGTGTTGAAATTAAAC
CCTGGAAGCCACAGAAGATGCCAAATGGACACATTTTGTGACTATTTATAGCTGATCCAAAAATGAAAGGGGAGAATGA
GGAAACTGATTGATAGTCCCCAATTCTGTTTTGATCTGAAATAGCTTGTTAAATGAATCTCATTGTCAGAGCTAGTTTA
AATAATAAATGTCTCTACTCTCCACTATTCCGTGCATATGTGAGTCTGAAAGTAAAAGTTTGTAATTAAAGTGACTGCA
TTTTAAAGTTTAAAAATTGATCTCAACTAATAGCTAATACTAATAAAATAAATAACTGCATTCTAATTTTTTAATTACC
CTGGTTTGGAAGCAGAAACATGTTTTGTACATAGCTGTAATGTTATTTCACTTTAACCTGGTTGTCCCCAGTTTAGTAT
CTACGAAGCTTTGCACGTATCGGAACTCCATACCTGCTTCTCCCTTGGTTTCTCTTCTCAGTGATAAAGGATTTGACTT
CTTTCTTCCCCATTCCAGCTTGCACATTCCTGAAGTTTTCCGGCTTGGAACAAATTATATTACTAGGTCTGCAGTATCA
TTTAGATTGAGAAGTATTGTGATAGAGCAAGGTTCACCCTAAGTATTGACTCTTAGAATCTGCTCATTCCATAGTCTGT
TGAGTGCCTTTACCTGTCACTGTTGTCCTAAGCACCAGGAGTGTAGCGATGAAGGATTTCTAGTCCTAGTTATCTAAGG
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Fig. 6. 190

ACTCACAATTAGGTTTCTATTCTGGGGCTATTTTGTACATTTGTGTACAAAAGACTGCATTTTTAAGGTGTCTGACTTT
TGACTTGGTTTCAGAAGGCATAAGAAGTTGGTTTCATGGTTACAACCCCATGCCCAGAAAAGTTAAGAGTGGTATTTTT
AATTAATTTGTAACTAGAATTAAATATAGCTAGAATTATATAGGAATACAACAAAAGATTAAGATGTATATACATCCCT
CTGAATAATATAAATATTGACAGGCTGATATTGGGTTTTCTAGACTTACTTTCGTAGGTATTTTCTTAGTATGAAGATA
GTTAAAATATACTAAGTGAAAAAAAAATACTCAAGTCACTGTATTAGTGGTTCCACTAGTAAAACTACAGACCTCATCT
GTTTCTGGAATTTGCAATTTTAAATATTGATTTTTGGAATGTGGCGACTTACACTGCCTACTTTTAGTTTGCCAGGTGA
GCTCATTTTTGAAAATATCCTATACTCTGCATTTATCCTTTGATAAAATCATGGCTATTATAAATTTTAAAAATGGTAA
AGATCATGTCAGAATAATGAATCGTGCTTCTAATTTGATAAATGTAGCTTAATCACAAATATACGAAACTTTCTGTTGG
TCTGGTGCAAACTTTCTCAAGGAGAAGATTTAGTGTTTCAGAACCATTGGAGTAGATGCTATATTAAACATTGAGGCTC
TCTCACCTTTAACTTTATAAAGCTAAATATAGTCAACAAATGAATTGGAAAGACATATTAGTAAAATCTACCGAGTCTA
GGCAGGCCTATGTGTCAGGCCAAGATGCTTCCTGGAATTATTCTGTTCAGAATGATTCCTCTTTCTTTCTCATTCTCTA
AAGATAGGATGTCAACAGTTTCTTCATAAATGTAAGGTTATAAACCATGACATTTGTGAAACTATTGCAATAATCTTGG
GTTGTGTAGGGATAGGATGACATTTTCAGAGGAAATTCTCTAAACACTTTCACATTAATAACATCTGGGAGCCATAATA
TTATAAGCATAAGAAGTGGATCTCATATAAAAACATAAAATATTTTTTTCACATTTATATGAAATAACATATGTACTTG
TAACAACTGAGAGTCCCTGATCAAATTAGGAAATCTACAGCCATTTATTTTAAATCTTGCTCTACCAAAATTGGATTGA
GAGACAAATTAAACACAAATCATTAGTTACATTTTATAACTATAGAGGATGTGCAGAAAACAAATATTCAAACCCAATT
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GCTACCACATGGAGAAAGAGCAGCTGGCTTTGCCCCACCATATTCAGTTGAAAGCTGCTTATTACTGAGCATTTGAATA
ACTTGAAAGGAGAGAAAAGGAAAGAACAACGTCAGGGTAGAATGTTTACTTGTTTGAGCCTCCCAGGAGAAATGCCATA
CAAAGATTTGACTTGATTTTTAAATCAAACCAAAGATCTTTCAGATTACATTGGAGGGCTGAATTGAAAGTTAACATTG
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AGAGTCGGGATAGATCAGAGCTCAAGTCTTAATGTCATCTCTTCCAAACTGTGTGACCTGGGAAAGACATTTAACGTGT
CTAAGCCTCAATTTCCTCATCTATCAAATGGGGATAATATGTTTACTGACTACAAATTAAGTAAAATAAGACCTACAAA
ACTGGGTGCTACTGTATGATAACACAACAGTTACTACAGATGCTATTATTATTAGGTCACGTGTGTGAAGAGAAAGTAA
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TGCTGTTCCTTCTTATTTCTCTTCCTTCCAATAATAATAATTCCTATTGTCTACCAAGTATTGAGAGCCTGTCATAGGT
CAGACACATGGCATGCATTGAGAGAAAGAGAATTCAAATTATTCTCTTGAACTTTTGAGCAACTCTATCAATGGGTCTT
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ACCAGAAATTTTTACTCCTGTGGAAAGAGTAATTGGATGTTCTCATTACAATATTTTTTTCTTCATGAATGTAATCAAA
CATCAGCTGGAAGCTTGCTTTTAATGCGTGGATAAACCAGCCCACACATTTTATTTTCCCTGAGGCGTATTGCTGGGAA
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CAAAAATTATGTTGTATTTTCCACTTCTAACATTGTTGATAGAAAAGTTGAGGTAAACACTTTATTCTGCCTTAGATGC
CTGCTAAATTAGTCTCTCTAAGTAAAAATGTCAGGAGGCTATTGTTGGATTAATTTAGGTGGAGTTGATACAGTTTTAG
GGAAGAACCGTTCTACAGTATGAGAACAACTTTTAATATTATGTTAGTTATAATCTGGAGCATTTATCATACCTTATTA
TATATCGTATGTTTTGACTTTCTCCTCTATTGGGCTGTGGGCTCAATTAAGATAAGAATGTATCTCCCTTATCTTTGTA
TATCGGTGCTTAGTGCATTGCCTGGCACATAGCAGGCGCTCAACTCTGTTGAATTAAATGATCAATGAGTTAAACTAGA
TGGCTATATAGTTGATCTTCTATATAGTTGATCTTCTATATTCAAAATATAGTTTACATATATGTATGCAAGTGTATAT
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AGCTAAAGTTGTATCAATATTAAAATTTATAACTAAAACGATATTACGTTGGCACATGAGGGTTTTAGTCAGCTCCAGCT
GCTACAACAAAATACTATAATCTGGATAACTTACACAACAACAGATTAATTTTCTCACAATTCTGGAGGCTGAAAGTTC
TGAAATTAGGGTGCCAGCATGTTTAGGTTCTATTGGGGGCCCTCTTCCTGGCTTGCAGTCAGTGCCTTCTGACATGGTG
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TCACCTCCCAAAAGACCCATCTCCGAATACTATTACATAAAGGGTTAGGGCTTCCACAAGTGAATTTTGAAGGGGACAT
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AAAATGTTTCACATAATTAGTTATGTTCACATGATCTAATGCCTGAAACTCATTTTTAAATATGCCTGAAACTCATTTG
TTAAAATTATAGTGTTGGGAAAAGAAGCATTTAATATTTGTTAGCATTGAACTTAAAAAAAAATTAAACACATTGACTTT
AGAGAAATGTATATGTTGAAGAGTAGGTGTGGATGAAATGAAAGAAGAGAGCTAACCCTTCTCAGTGCCAGACACAGTG
GTGTGGGCATTAATAATTCAACCCTCATAGCAGTCCTTAGGGTGGTCTTAACCATTTCATGGGGAAATTGAGACTCAGA
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TTAGGGACCTTATCTGTTATTTATATGAAAAAAAACAACTAACTTTAATGTTTATGGAGTTTGTTTAATTAAATAAAGC
CCTTAGGGATTTCAAATTGTTTCCTAAGTCCAAGAGATCCCAATGAGAAGGAACACAAAGTAATTTTTAATGATTTAAAG
TAAATGCTTTAGACTGAAATGGTAGCTTCTTTTTAAATAACATTTCAAATTAGGTTTACTATTAACATTTACAAAGAAC
CAGGGCTATAAATCAACTTTATATGCAGTGCATTCCATTGATCTTAATAGTTGTATTTGCAAGCTGAGTAGAAGAGATT
TACAGTTCTCTCTCAAGGGATAGATCTCCTTTTTTGGGCAATGAATATACTCCTTCAGAAAGTCCTCTTGCTCTTTTAC
GTACTCTTTTTTGCTTGGTATTTCAAAGGCTTTTTGTTATACTTTAAAAATATTCAACAAAAAGTCACTGAATATCTAC
AATGTGTTTATATCTTGGGTGTAATGCCTAGGTTAATAAGATGCATAGAAATATGGACCCAGTGCTACAGGCCCAGCAG
TGAGCAAAACTCAGGGAAACAAGTGACACTAAACAGTGTGGAGAATGCAAAGGCTAAAGGCTATATGAGTCCTGGCATG

Fig. 6.191

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AGCACGGAGGTATGGGCTGAGCAGCGCTGCCTGAGGAGGTGATGTACAAGTGGGTATTTATCCCCCCCGGGCAAACCAG
AAGGAAAGGTAAAGCCAAGGAACAAAGTAGACAATGGAGATGGTGGGGAATGGAGGGAGAAGAGGAAAGAATCTTGGAG
AGATAGTCAGTCTTAAGGTCCAATTAAACTATTTCATGGCAGCCATTATTATCCTGAATTTTATGGATGAAGAAGTGAG
GTTCAAAAATATTACATAACTTGCCCTGTGTGACAAAATAAATGAAAAGCAGAGCCAGGTTCAATGAAAGTATGTTTGA
CTTAGGTTTTAAAATTCTAGTCATAAATGGCAATTTAATGTTGTTATATATTTGTATTGGGCTTCAGCAAAAACAAAAA
TAAAACCTCAAGTACACAAAAGCTCTAGTAATAGAGCCATGTTTGTGCAGTTATTTCCAGCAATCCTTGGAACCTCCAA
AATTCTCCTCAGCCTGACATTATATGCCCTATGTGTCATTTCATTATGTCACCACCTGGTGGTGCACTCACACCCACTT
TGGGAGAAAGGGGGTTAGAGAAGGGAGGACATCTGTGGACAAGCCAATAAAAGCATTTACTGGCATCGTGACTTCATAA
TTTACACTTTTATTTATGCATAATTAAGTATAAATAAACAAAAACATGTATTTTCTAACTGCTACCTGTTCTGTTTCCA
GTCTTGCTTGAAAATCATCTTTCTCAAAAAACTACCTATCATGTGCTATGCCCATTACCTGGGTGACAAAATGATCTGT
ACACCCCTACAAAATGCAACTTAGTCATGTAATAAAGCTGCTTATGTTCCCTCTAAACCTAAAATAAAGGTTGGAAAGG
AAAAAATAAATAAGATAAAAATTATCTTTCTCTATCAGAGTAATTTGACATCTTGAGGAAGTGATCCTGGGACTTCATA
TTCTTTAGGATTCAGGTGTCCAGATAATCCCAGAAGTAGCCAGCAATTTGGCCATTTGGGGTGTAGAAACCTTCATACC
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TCTAGAAGAGTCTTCAAATAGGTCCTAGAAGGACCAAGTAAAATCACCATCCCTCAATCCCTCAATTTTTCCTTTTTCC
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TTGGGATAAAATTGTTTAATGTAATTAACTACAAGGAGAAAAGTTAACCAGTGGCTTCTGCTTTTGCTGAAAGCACTTT
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TAAAGTATAATGCCCAAATAGGCCCTTGCCATTCCTATCAGGGACATTGCATCCATAATCCATTTTCCATATCCGTTTG
TATGTAAATGAAAAGTCTCACACATCACAACTTCTGTTTTTCTCATTGTAGGATCGCCTCATCTGTATTTATCCACATT
AGTAAAAATTATCCAAATCCCAGAAATATAACAATAGTCCGTATTTCTTGAGCATTTACTCTGTTCTAGGCTCTGTGTT
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AAAACATGTTGAGTTATTCAGGTAATTTGCCTATATTCACATACTGGTCATGAGAAAACTAGAGGACCCAGGAGATTGA
AGCCCATATTTCATCATTTGACTGTCCTAGGAAACGCCTTGTATACTTGTTTTTTTTAAGGTATTCTGTCACCCAAGAAC
ATTGAAGGTATATGCAGATTCTCTTTTCCTGTTCTATTCCACTAGACCTGAATATGAGGGATGAAAATTGCTTTGGTTT
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TATTAGTTTATAAAGAAGGCTGAACCATGACTATATAATAATGAAGCAATTGTAAAAAAATCAGAAAGCATTCACATTC
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GACAATGAAGATTTGCAAAGGTTCTAGAATACTTTAAATTAAGTTAGAATAAACTTTTAGTTGCACTGTGCTTTGACTT
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AGAGACTTGCACATTTCTCACTCATTAATATGATCAAAAGCATCTTATTTATGTCAAAAGACGAGAAAACGAAGGCCATT
TTTGGCATTGTCCTAAAGTTGTGCTTTCTTGCCTTATTAGCATTTCATTTCAAGCAATAGATTGAACAGAAATGCTTGT
GATTGAATGAAGATTAATTATAGGGTATGGGAATATTCAAACCTTTTTAATTGTTCTGAGTAGTGTCTTCTGCTGTTTT
GTTATCCAAAAGGGAGTAAGTATTTGGGGAACAAAGATTGTGACACATCTGGTAATATTCAAGATGCACACCCCCTCAC
TAGACTGTCAAAAGGCTGGGCTTGTCACAGATGTCGAGGCTGTGATGTATTGTCACTCTTGCTGCACCCATGGATGCGC
CTCTCCTGATATGCGACTCCCATTGGGAATGTGAAGACACAGAGGAGATTTTTAATCAAGAGCATAGGTAGGAGTCAAT
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CTTTTATGCAGGACTCTGACCTAATTATTTCAAAAATTATTGCTACTGATTATAATCTCATTGTTGCTTTCTTTTATCT
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GACCCAGCCTACCTTGGAAAATGATTAAGCCCCAATTTTCCTATATGTAAAATGGAGACAATACTAGCATCTACCTCAC
AATATTGTTGATTGAATGAAATGAGATAATATAAGTAAATTACTTCCAGTAGTCCCTAGCACATAAGCACTCATTAAAT
GTTAGCTTTTAAAATTGTATTCCAGATTAACATGCCTAAAAACTAGGGCTACGTACTGAGAATTCATAGAACCAATTTT
ACTTTGTGTCTGTGTTCTTATGCCTGCATCAGGCATTACGAAGAATACAGAGAGACAAACTTTCTATTCCTCAGGGTTG
ACAATAGAAGAGAAATAATAATACACAGGAAATAAGTAAAAGTATAATTCAAACTCCAAGTTTAATGATGTATAAAATG
AAGAGGTCAGTTTGGGCTATAACAATCAGATAAGATTCACTAGGTGCTAAACTCGATCTAAACTGAGAAAAATAGACAA
GACTTATAAATATTAGTTTAAAACATAAATCTGAAAGTGTTACTCCCCTATATTTAGTAAGACTTCTCTGTGGCCCTT
GCTCCTATAACAGTTTCTTTAAGGTTAAAACTGCTGGGTTTCTGAGCTCTTCTCCATCCTACTAAATGAGAACCTCTAA
TACAAGAGGACAGTGCTTGGAAATCTACATTTGTATAGAACCCAAGTGATTCTTATCAAGCAAATAATGGAAATTCTAT
CTTAGAGAAGTCTAAACATACTAAGGCCCTTGTTTAGACTCTATGTAATTTCATTTTCAGCCACTCTGGGCCTTGACCC
CTGAGAACCAGCAAAAGAGAACTTTCTTCAGCACTTGGCTTAGCCCTCCCCTCTTTTGTGAAGGAGTTTCCCTTCACTG
GTCAGTGTTCCTTGTCTTTAATGATGGTCTTGAACACCACTATGTTGTTGTACTTAAAACACAGAGTTGTAACTGTTTA
CAGGTTTATATCCATTAAACCACTTGATCTTGTGTTTCATGCTCTCTTATTCTCACCTTAGCATTTAGTTACCACTCAC
TAAATCAGTTGATCTTGAAACTTTAGAAATCTTACCCTTACCTGAGGGGTCCTTGTTTACAAAGCAGATTTCTAGATGC
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Fig. 6.192

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TCTTGCTCAGGTACTCTTATTCTGTAGTTGAAATGGGCATTTTAGGGCCAGACACTTTGGCTCATACCTATAATACCAA
CACTTTGGGAAGCTAAGGCAGGAAGATCACTTGAGTCCAGGAGTTCAAGACAGGCCTGGGCAGCATAGCAAGACCCTCT
CTCTACATAATTAGCCTGGCATGATAGCACATACCTGGCTACTCAGGAGACTGAGGTGGGAGGATTGCTTGAACCCAGC
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TCTTGAGAAATAGCAGAGTTTGAATTGAATGTGGCTTTTTGCATAAAGTGGTAATTTTGGAATTTTGGAATAGGAGAAT
GAATAAAGATGGCATCCCCAATAGTTGTCATGGCTAGTCTTAGGAAGGGTTGGATCAGATGATCTCTAAAATATCTCTT
AACACTGAAATGTATAGTGTTATGCTATAAAATATGTAGATATTGAGTTGTATTAATAAAGCCTTAGCTTGTATTCATG
GGATAAAACTTCTACTTAAATAATATTGTAATTTCATTCATTAATGCATGTGTATAAAAGTGGATCAGTATTGTGGAAG
GTTTGCTTAAGGTGAGATACATGCAATGGTAGCCTGGAGCCAGGTAGTAGTACCAGCTCAGGAGTACTAACTGTTACGT
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AATGCAAAGGTAATCAATACTCAAAACTCATTAGTTCCCAAGTACTTCATTATATTTTACTATTATCCATGCTCTTGAG
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TCAGGGCCATCACATTGGTAGCTTGAAATAAGACATGCTGATAGTAGTTACACCATGGAAATGGACATACTGCACAAAT
CAGGCCTTTTTTTTTCTCCTGGAGAGCCAACTGATAAATATTTACCAGCATACCAATGGCTCATGTTTAGAATAGTCCC
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GAAACTTCTGAATTTTTCATGTATGCCTTCAAGAGTTGTAATCCTTACTTAAGCAAATGTCTGTTTAGGAGAGATTCCA
GAGATTTTCCTATTTAGAGTGTTTTAAAAATTAAGGAAGGTGTTCACCAGCTTACTAGAGGTTCCTTGTGAAAAGAGAA
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GTGAGTTGTAGTATGAGATGGTAATTTACATGGGGTGCTCCATGTGCTTAGCTTTTCTCTGAAGTTATGCATCCCCACA
GCAAGGGAAATATTTGCATTTCTGAGAGTGAGAATTTATGTTCATCCTCTACTAGGAATGGTGGCAGCTTTTCCAGGTC
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AAGATGTATGTACAGCGTCAGCAGCGTAGTTGTACCTTTTACAGACAATAGTGGCTCCAAGCCAAGCACAAGCTTACAT
GGGTGATCACCTAATGTGCCTCACGTGGCGTGGTTACATAATGTGCAGAGTTGTGAGCCTGTGCTCCAAACTCACTGAG
TCATGCTGGACCGGATGTCTGCTTCGGCCTATTTTTGAGAGCAGCACATCCATTTTCCTTACACTCCACCCTCAATGCC
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CACAGGTCTGACACATAGGCCTTAAATTCTACTCCCTAGGCTGAAGGAGTCTTTTAAATGGAGAAACATGCCCACAGGG
TGGAACCCCAGACCCAGAGGCCACAGCAGTAATACAAGGAGCAACAACTCCAGGTTATGACGGGCAAACACCCCATGAT
GATGTTACCCGAATTTACTTTATGCATTAAGCCAGGTTTTTATTTCCCTACCTTTAGGGGCATTGGGGCATGCAACAAC
```

Fig. 6. 193

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AGGTTACCATCCGTTTCCATCCTGGTCATAGGAGGTCATCCTTCCTCCCATAGGTCTTGCCACATGGCCTAGCCCCACA
TGGGCCAGTGGCCAACTAGCCACTTCTGTATTTTCCAGGTGCCTAACCACAAGGTTAAGCC'LJATAGACCACACAGCT
ATTGGTGCAGATTATCATATGTGTCACCTCTTTGGTGATCACCATTCATATTGCTCTGACTTCAGCCCATTGACTACTT
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TAAGGGATTATACTAAGGGCAATTATGAGAACTCTTGGGACAAAATAATGGTCTTATATCAACAAAAATTCCCTGGAAG
TGᵀGATATGTCAGCTTGCAGAAAATTTTCCCAATGTAGTGATAAGAATTCTCTGAAGACATAGGCAGATTGAAAAGACA
TGAᴵTAAGAAACAATGTGAGATACATTTCCATACTTACTGCAGAGGTAAAGCAGAATGGAATTTGTATCTTTTCCTCTG
CAᴳGTCCTATTAGAGAAGACAATAGAGAACCCTCTAAGAGGTGATTGTCAAATTGGTTAATATGCTGTGGTTTGGAGTG

Fig. 6. 194

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CCTGCATTGTGAACAAGGGCACATTCATTCAACTTTCCACATCTTCCTTTCCCTGTAAAATGTACATGGAGTTATGGTA
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AGGTGGTAGCTATTGGTTAACCCTATTCTTTATAACATAATTGAAATTAGGTTATATGACTTTTAAATATCCAAACAGT
CAGAATAGAAATATTGATGAGAAATTTTGCTTCATCTCAAAAACAGTGTTAGAACTACACCAATGGAGTATTTTTCTCC
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TTCTTACTATTTTATTTTTAACATTGACATTAAGTTAAATAATATATTTTGTTACTGTAGCCTAGGATTTTATTTTCCC
TGTTATGAGAACAAGGATAATTATTTTACAAATGAAATTTTAAGGGATTAGAGTGCCATAGACATGCTCATTGCATTAT
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GTTCTTAGGAATTTTATGCTTAGCTGAAAGTATTAAACACTTTTCCTTATTTCCTCAACATGAATTCTTTTTCCCGGGG
CTGCAGGGGAAAGGGCTCTGATCCTGGTTCAGTCTCTAGCTTGATCATTTTAGTCGAGTTCCTTTACTTCCATGCAACT
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GATGTGSATACCATTTTCTTTAGGTTTTAAGAAAACCACTCTAATATTTTTTCAGGGCCATTATTTTATCAAATATGTA
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AGATTTGTTTCAGCCCTTCGATTTAATCACATCATGTCTGATATCCTAGCATTTAATTACCCAAATTCCCAGTGTTGTT
CAAGAAGGTAGGTCTGTGGTGGGTTAGCCTTAAGCTAGTGGCTGAAAAGGTCTTGATTTGCCCCCACATCTAACCTAGT
CACTTCCTGAGCTGAAACTCCCATATCACCTGGGGACAAGCGACCCTGAAATCCTCACTGGCCACAGCTTCTTCTGAGG
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Fig. 6.195

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GAAGCTGCAGGAAAAAGAACAAAAATGAGCTTTTGTGAAAGGATTAACTTGGGTGAAGAACATCCTCCATAACCAAATA
AATGGAACCTAGTAACCCACATGGCCAATCCCCAGGGCACTAGTGTCTTTGGGGATATCAGGGTGTGTATAGAATTTCG
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CCCAACTTCCTGCAGCTAGGATTCAAATACAGACCATCTAACTCCAGAGCACTGCCCTTTAACATTGTGCTATAATCCA
CCACTTTGCATAGTTGATCCTACAAGAGGATAAGTAGAGATCAAAGTAAGAAGTGATCAATGGATTACTATGTCAAGCA
TGATTCCCTCAAACTATTGGGAAGAATAAATACCATAACATGTATAGGGTGCTGTGTACAATTCCTGGCACAAAGTTAA
GTACATGATAAAATGGAACTATTAATATTAGTAACCCAGATAGGTTCTGGGCACATTAGCAGGAAAGGTCTTCTTAATT
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TTTCACTAATTACAATAAATGTTGAAAGAAGGAGTTTCACCTAGTATCCTTACTAATATCAGGGAATGGCCTGGGCTTT
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TGAATGTATGTATGTTGCTTCAAGTCATTCATGCACACATTGAAGAGGACAGACAGAGCCCTGCTGAACACCACTAGCA
AATCTCTCTAGTAGACCACAACAAGTCAATCTGTGCTCTTTGATTACAGTTGTTCAACCAGTTACTAATCTAACTAA
TTGTAACTGGTTGAACAACTGAACCCAAAGGGTGCAAAGTAGAAACATTTCATTGTGAAGTCGGCTGGGGACAAAGAAA
ACACCACCAGAACCCCCTTCAATATGCAAAGGGAAAATTCCTTCTCAAGTACAAGCTACATGGTGCCTTTTTGAATTAT
CATCAATTTGTAATGCTGTATCAATACATATGTATTATCTGTTCTTTTGGTAAGGGTTGTTTGAAGTATAAGGTACTTC
CTCCATTAAGGTGATACTAACCACATGTTAATTATTGTCTGAAAGTCACCAAGGATATGAATAATAAAAAGTTTTAAAA
ATAAGACCTGTTTCTTTTATTGATAGTTGTACCTCAAGGCTGAAGTGAGGTTTGCAATGTAAGTTGTAAAATGTGATGT
GAAATAGACAATTCTTTGTAGTTTATATAAGGCAATATATCCATGTGCAATTATGGTCAAAAACAGCAGACTTTTAAGT
GATTATTCTAAAGTTATTTTTTTCCAAAATAACTTTATTACTCTTGATATCATACCATATTCACAAAACATTCTGGTAA
AGCTATTGCTCAGTGTGTTGTCCCAGTGAGACTCAGGGAACATTTCTATGTGACGCTTTAGGATTGAAGACAGTTCCAC
GTTTTCTGAGTAATTCCAAACTGTGTAAGAGATTATGTTCCCTTTGCATATTGGCTGCTAAGAAGCTCACTTTTTCACT
TATGATACTTGGATGCATTTTGGCTTTTTGGGTTTTTGTTTTTTGGTCTAATTTTTTTTTTTTTTTTTTTTTTTTGA
GACAGGGTCTCACTCTGTTGCCCAGGCTGGAGTGCAGTGGCACCATCATGGCTCACTGCAACCTTGACCTCTCCTGGCT
CAGGTGATCCTCCCACCTCAGCCTCTCAAGTAGCTGGAATTACAGGCATGTGCCACAACACCCAGCTAATTTTTGTATT
TTTTGTAGAGATGAGGTTTCGCCATGTTGCCCAGGCTGGTCACAAATTCCTGGACTCAAGCCATCTTCCTGCCTTGGCC
TCCCAAAGTGCGGGCATTACAGGTATTAGAGGTAGGAGCTACTGCATCTGGCCATGGGAAACATTTTGAATGATAACTT
TGTTTGTTTGTTGTCTGTTGAGCAGGCTGGAGTGCAGTTGCATGATCACGGCTCACTGAACTCTTTCCTTTAATTTTTT
TATTCTCTCAGCACGATTATAATTTATTTTAAAAGTAGGGAATAATTGGCCAGGCGCAGTGGCTCACACCTGTAATCCC
AGCACTTTGGGAGGCYGAGGCAGGCAGATCACAAGGTCAGGAGATCGAGACCATCCTGGCTAATATGGTGTGAAACCCC
GTCTCTACTAAAAAATACAAAAAATTAGCCGGGTGTGGTGGTGGGCACCTGTAGTCCCAGCTGCTTGAGAGGCTGAGGC
AGGAGAATGACGTGAACCTGGGAGGGGGAGCTTGCAGTGAGCCCAGATTGTGCCATTGCACTCCAGCCTGGGCGACAGA
GTGAGACTTCATCTCAAAAAAAGAAAAGCAGGGAATAATTAGTAAGTGCCTAACTAATCAAAAATTGTTTCCAGGTGAT
GTTGCTGTTGATGTATGAAAACCTCTTTCACCTAGCTTCCCTCTTCATTCTTTTCTGTCATATTTCTATTGACCAGTGC
TTTCTTGGCCCTTGGAAATGTGATTAACTTTTGCCATGACCTCTATGTTAGTGCCACACCTGACACCTTTATGCCACTC
TCTGCCTCCAGAACTCTGTGTCCATCTTATATGTTTTTACCTTCTGCTGGGCCTCAGGTCTGGATGCAAAGCTGTAGAG
GAATTGAGCTGTTTTTCACAAGGGATCCAAGAACACATAGCATGTGTGAACTGTACTAAAGCTTTGAGAAGTTGTGAAG
AAGATTCACATCAACTATGCACCACACAGCCTTCTTGTCTTGCTGGTACCATCTCCTCACCTCTGTGTTGCTCCTCTTG
GAGTCTTCACCTTCCACAACCAGCACTTCTATTCTCACTGTCTGAGTGTGACTTTCCATAATTGTGAAACTCCTTGGGC
TTATCCCTCACACTCATTCTTGAATTCTAAGGAATGGCATTTTCTCCATTGTGCTTACATTTGCTTTTCAGTATTAATT
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Fig. 6.196

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GTTCTAGAGAAAAGCATTCATTCACTCACTCACTCATTCACTCATTTAACAAAACCTGTATGGGTATCTACAGTACTAT
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AATGAATGTAGGGTGGAGAGAGGAATRGGAAGAGCATTTCAGGGAGAGAGGAATAGCAGGAACAGAGACACTGATGCAA
GGAATCCCATGGTATTTGTGATGAAGGGATGATTTAGTAGTTCAGTGACAAGACATGAGCCTGGAGATAAATAAGGTTT
AGGTTCTGGGGACCTTGTGAGCCAGGTTAAGGACTTTGGGTTCATCCTCAAACCATGAAGCATTCCAAGCAATTATGTG
ACAAGATCAGATTCTAGGGGCTGAATGGAGAATAAATATGTAAGTGGCAAGATTGAAGGCTGGGAATGTAAAAGGTGGC
TCCTGCTGTGGCTTGGGCAAAACATCACATAGGCTTGAGCTAGCTTTGTGATTCTACGGGATTGAGAGGAGAATGCTAA
TTTCATAAATGTTTAGTGGATATGATTGGCATGTGAWCTGAATGCAACTGGAGAAGGGGACCAGTCATTTACTGAGTWG
ATATTGGTAATTGATAAATAAGACCAAAGAAGGTGAGGTTTGGGGATAGGCAGTGTAAATTTTGGAAGTAATAAGGGGA
TTGAGGTTCTGTCCATGGCATTGGTGGGGAAATCTAGCATGTAAATACATGTAAATAGATAAGTAAATATGCCTATACA
TTTATATAAGATATCTATACACTTATTCTCCTTCTTTGTTCATTTATTAGGTGTGAAATCTACGTGATTTTTTTCTCCC
ATATTATTTTTGGTATTCAATAAATGTGAATCATGAAGAGATAGCTGTTTTATTTAATCTCTGCAAAGTCACATTCTTA
CCCATTTTGATTGTAATAAAAGGCCTATGTCGTTATTTAATTTTTTTAACCTTTGTGGCATGATTTATAGAAGGAAAATA
AAAGTGTCATAGGTATGATAGGCCAAAGTTGGGTTGTTGGTAAAATAATAATACCACAAAATGTTTTTCTATTTAGCCA
ACTACTGCAAAGTTACTTTAATTGTGTTTGCAATTAATTGATTTCACTTTAAACCAGAAACAATAGAGACTGTAATCAC
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TGCAGTCACCATTACCTCCACCACCATCATAATTATATATAGCTACCACTTCTTGGACACTCAGGTCTGCTTGTGTTTG
AACCAGACATTTCTAGCTCCAAAGCCTGCATTCTTAATTTCCTGCAGTCTTAATACCATCTTTTTAATAGAAGACCCCC
ACAGGAAAAGCTGATTCTATAATTTAAAAATGATTTGGGGATCCAATAAGTCATGATTCTATTTTATATAATTTTGGAA
GATGTGGCCAGAGAGCTAACTCTTTTTATTTTATTATTTTTCTTTTAATTGTTGACGTACTTATAACCACCAGCTATGA
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GGAAATTAGGCAACTGCTAAGGTGGTTTTGCAGTTTTCAGAGACACAGATTAGGTTTTTCAGCCTTATATAGATCTG
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TGTTGATTCACTGCTGTCATCCCTGTCAAGAGAGCTGCATATGTAATTATGGTTTTTCTTYTGAATAGATTTGCTTTGGG
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CCGGGGTCAAGATGATTACTATGTGATCTGCTTAAGTGGGAACAGGGCCATTTCCCTCGCCTTCAACACACTCAGATAC
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TGCTGAAACTGATGTCTATAGAGCTTCAGTATTTGTCCACTGCATAAAATCAGTAAATAAAGGCAGGAAGTTTGTTAC
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GCCCTGACCTGAATCAACTCTGAGCCCCTACCTAGGCTGTTGTTTCTTAGCAAGTACATAATAACAACTCATCAGAATA
GCAGAGATCTTGGTAACAGGTTGCCTGACACTTGACTGATATGGGAAAATCCTGAACAGGTTTGGACTGGAATCTTGGC
TCACAACCTGTGTGACCTTGGTCAAGATATTTTATCTCTCTGAGCCTCAATTTTCTCTTCTGTTAAATGAGCATAGTGA
AACTTTGTTCATTGGTTTGCTTTGAAGAGTAAATGAGATGGTAAGTGTTGGTGTATGTAGTGCAGGAACATAACAATTC
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GCCCTTCAGGTCTCAGGTTCTCCTTCCTGGTACCCCCAGAAAGATTAAGCCCCTTTGGCCCTCTATGTGCCTCTTTTAT
AGCATTTAACACAATTATAATTAATTGATTATGTAACTTTTTTATTTAATAGCTGTCTCTTGTAGAATATAAGCACATA
AATTCAGGGACTATCTATTAATGAATTTTCCTCTAGTGCCTAGTACTTAGGTGGCTCACAGTAAATGCTGATTCAATGG
ATGAATAACTTTATTCAGTCTGCTCAGTGTGGCCTTTAATTTAGATTCACCCCAAATTACTCCCCATCAGGAAGGCTTT
CTTATTCAACTTTCTCATCCTCAAGATTAGCAAAACTACTCTCTTTCTATTCCCAAATATCTTCTCTCAGTTTCATTAC
CATGATCAGAAAGTACTCTTCCTTGAATTTATTTTCTAGACTAGATCTATTTTGGGGTTTTCTTACCAGTGATTGTAAC
TATTGATGTGGTCTTTTTTCTGTTTTTTACTTCTCATATGTGTATCTCTTTCCATTAGTCCTTCATTCCTTCAACAGAT
ACTTTCTGTTGTGACAGCCACTCTTCTAAGCACTATGGAAGGATTTAAAGCTGTGTGAAGCACAGTTATTCCGGTCAAG
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Fig. 6.[192]

GAACTTAAAAAAATAATACAAGTGACAAAACATGTGCAAGTTTGACTAAAGTATGAGACTGGAATAAATACCTTATAAA
AACCAAGAGCTTCAGTATTTTCCTGCAGTTTTTTCCTCTATTGTATTCTCAACTTTTACTGGTTTCTCATCAAATAAAT
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TTTTTCATGCTGTATAATGTATATTAAGATATTGCTTATTGTCTGTCTCCCTTCAATAGAATGTAAGCTCCATGTGGGC
CAGGATGTGTAGTCATTGATGTATCCCTAGAATAAGTGCCCAATTATAGTAGTCATTTGATGGACATTTATTAAATGAA
ATTATGAATGAAGTAGGGATTGGCAGTTATACACTAAGACTCTGGAAAAATGGCCTGAGTTGAATCCTGGCTTTACCAT
TCTTGGCTATATAATCTCATATTATTTACTTTCAGCCTTTCATTTCCCAGATGCCCCTACCTTCTAGGGTTGTTGGGAG
GATGGAATGCATAATACATGTAAACCAGCAAGTTCATAGTAACAAAAGTTGACTTTTTAAAAAGTTAACTTACTCTTCT
TTCCTATATATGTGTAAATCATATTTTATTTTCTCATTTTAAAAAGAAGACAATAACTAAAGGTATTAGCACATAATGA
TTCAAAACATAATATTTCCACATGTCGAGTTGAATGACTTTGAGCCAGATAATATGAGTTGAAATCAGTTTAACAATAA
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TGGTGGCTCATGTCTGTAGTCCCAGCTACTTGGGGGGCTGAGGCAGGAGGATCACTTGAGCCCAGGAGGTCAAGGCTGC
AGCGAACTGTGTTTGTACCACTGCACTTCAGCCTGGACGACACAGACCATGTCTCAAAAAGAAAAAAAATAGAATAAAA
TCAAATACACACCATTTTTGAATATGTCACCAGTCTGTGTAGTTCATCTTGAAAGGACTTCAAGGTCCAATATGCATCA
TGCAGAGAGTTGCTAGGGCCCAGACAAGAGTGACTTGACCATTGGCCTGAGTAGTTAAACTATCAGATAACTAGTGAAA
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GTAATGTTTGGTAAAAGTCCTTTAGAAGTAGACTGTTATGTGTGTTACTAGTTATAATCAATTAATAACCTGTGATTTG
TAGGAGCAAATGGTCATAGGGATACAGTATACATTTTAATCTTGCTCTTCAAACATCACCGTAGATCCATGGTCCTTCT
CAAGACATTGGCTTTGTTCTGAAGCAGCTCCCACGCTCTTCCAGAAATCTCTATGCGGGACTCTGAATGTGGTCAAGAA
GAAGATGTACTGGATGCACATTCCCTATCAGGAGTCTCTTAATAGTCTCCCACCCAGTTACAACATATTGCTGTAATCC
CACACAACAGCTGAAACATCTTTTCTTCATTTCTTTTAATTCCTGTAGCATTTGATGTCTCCACCGTGTAATTTACATT
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CCTCCCCGAGTATAAATTCAAATGCTATTCATTTTCTGAGTTGCCTGTATTTCTTTAGCCCTTAAGGCATCAACCTTTG
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CAGAACCTTATTATCATCACAAGGAAGTAGATTAAAAAATACTTTTCAGTCATTCGTATTCAACAAGTACACTCCATCA
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ATACATTAAATAGACCTTAGCAAACAACTTCTTATTTGAATACTTGTCTGTCCCTGTCACTCTTCTCATACTTTAGCCT
CATAACTTTGGTTATAATTATGTAAATGGTTAATATTCATGTTCTCATTGCAAAATGAAAAGTGAGGAAGAGAAATTAA
GCCATTTGCCTAAGGTCACAAGTCTGGTAAAATCAACAGAGGCACTCAGAATACCTCCAAAATCATTTCCATGATGCCA
GAACCTTTAAATGCTACAGAAACAAGCTAAAGCGATGCATTTAAATGTGCTTCTATGTAGGGCTTGAGCTGTATCTAAA
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TAAAATGAGATTGCAGAGGAAACACATTTTAATACCTGAGGTGTGTGCTTAATCTTCTTGATGTATATTAAAAGCTCAG
TACGTGAGAGTAATATGAGGTGATGGGGTTTACTCTTAAAGAGATTACTAATAATGTTTATTTGGAAAAAGATGAAGAT
TTTAGAGGCTATTTAAGAAACTGGTTCTGGGAAAACCAGCCATAACTTAAGAGTTCTTCTTTCACCCAATCCCCTTGGA
ATTCTGCTTTGCTATAGAACCATGGTCAAACCAAGGCAAGCAGCACAAATTACATGAACAAAGAATACAACAATAAAAA
GCGGATCCTTAAGACCCTCCCAAGAAGACCAAGAGCTCTCTCTTTAAAACTCTGTAAAATACCTAGGTTCTCAAGTCTT
CCTATGTCTTATTTCTTAACTAATTGCACTATCAATTCAAAAATGGAACAAAGGATATTCTACGTATCAGAACCTTTTT
TCCTATACATTAAGAAGAACTTTTCCCACATGAATAGGTAATATCACAGTCTAAAGCCAGAGGATGAAACCTATGAATT

Fig. 6. 198

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CTTCCTATCATATATTTTAACAAGAAACGTAAATATCTATGACCTACTTATAGCCAATTTATATTTTGCCCAGGTTGTT
TTGTTTCTAAACTTACCCCTCATATGGCTTAATAATGAAGGCCATAAATGTGCCTCTTTCCTATCTCACCCCTATGACT
TGACATGTATTATAAAATGAACTTCCTATATAAAATAATATTTATTGAATGAAAGGAAAGTATTACATACAACTTTCAT
TATTATATTATTAAGCTTATTTTTTTGCACATCATRGCTAGATCATCTTAAAATAGTTTGGCTCTGTCTCTTTTCCCATT
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CATGACAGACCCCGGTGTGTGGTGTTCCCCACCCTGTGTCCAAGTGTTCTCATTGTTCAATTCCCACCTATGATTGAGA
ACATGCAGTGTTTGGTTTTCTGTCCTTGCAATAGTTTGCTCAGAATGATGGTTTCCAGCTTCATCCATGTCCCTACAAA
GGACATGAACTCATCATTTTTTATGGCTGCATAGTATTCCATGGTGTATATGTGCCACATTTTCTTAATCCAGTCTATC
ATTGATGGACATTTGGGTTGGTTCCAAGTCTTTGCTATTGTGAATAGTGCTGCAATAAACATACATGTGCATGTGTCTT
ATAGCAGCATGATTTATAATCCTTTGGGTATATACCCAGTAATGGGATGGCTGGGTCAAATAGTATTTCTAGTTCAAGA
TCCTTGAGGAATCRCCACACTGTCTTCCACAATGGTTGAACTAGTTTACAGTCCCACCAACAGTGTAAAACTGTTCCTA
TTTCTCCACATCCTCTCTAGCACCTGTTGTTTCCTTACTTTTTAATGATTGCCATTTTAACTGGTGTGAGATGATATCT
CATTGTGGTTTTGATTTGCATTTCTCTGATGGCCAAGTGATGATGAGCATTTTTTCATGTGTCTGTTGGCTGCATAAAT
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TTCCATTAGTCTATATCTCTGTTTTGGTACCAGTACCATGCTGTTTTGGTTACTGTAGCCTTGTAGTATAGTTTGAAGT
CAGGCAGCGTGATGCCTCCAGCTTTGTTCTTTTGGCTTAGGATTGTCTTGGCAATGCGGGCTCTTTTTTGGTTCCACAT
AAACTTTAAAGTAGTTTTTTTCCAATTCTGTGAAGAAAGTCATTGGTAGCTTGATGGGGATGGCACTGAATCTATAAAT
TACCTTGGGCAGTGTGGCCATTTTCATGATATTGATTCTTCCTATCTATGAGCATGGAATGTTCTTCCATTTGTTTGTG
TCCTCTTTTATTTCRTTGAGCAGTGGTTTGTAATTCTCCTTGAAGAGGTCCTTCACATCCCTTGTAAGTTGGATTCCTA
GGTGTTTTATTCTCTTTGAAGCAATTGTGAATTGGAGTTCACTTCKGATTTGGCTGTTTGTCTGTTATTGGTGTATAGG
AATGCTTGTGATTTTTGCACATTGATTTTGTATCCTGAGACTTTGCTGAAGTTGCTTATCAGATTAAGGAGATTTTGGG
CTGAGACGATGGGGGTTTTCTAAATATACAATCATGTCATCTGCAAACAGGGAGAATTTGACTTCCTCTTTTCCTAATTG
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TAACTTTCTGTCTCGTTGATCTATCCGATGTTGACAGTGGGGTGTTACAGTCTCTCTTTATTATTGTGTGGGAGTCTAA
GTCTCTTTGTAAGTCTCTAAGGGCTTGCTTTATGAATCTGGGTGCTCCTGTATTGGGTGCATATATATTTAGRATAGTT
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CTGTTTATCAGAGACTAGGATTGCAACCCCACCTTTTTTTTGTTTTCCATGTGCTTGGTAGATCTTCCTTCCTCCCTT
TATTTTGAGCCTGTGTGTGTCTCTGCACGTGAGATGGGTCTCCTGAATACAGCACAGTGGTGGGTCTTGACTCTTTATC
CAATTTGCCAGTTAGCATCTTTTAATTGGAGCATTTAGCCCATTTACATTTAAGGTTAATATTGTTATGTGTGAATTTG
ATCCTGTCATTATGATGTTAGCTGGTTATTTTGCTCATTAGTTGATGCAGTCTCTTCCTAGCATCGATGGTCTTTACAA
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GGTGGTGACAAAATCTCTCAGCATTTGCTTGTCTGTAAAGGATTTTATTTCTCCTTCACTTATGAAACTTAGTTTGGCT
GGATATGAAATTCTGGGTTGAAAATTCTTTAAGAATGTTGAATATTGGCCCCCACTCTCTTCTGGTTTGTAGAGTTTCT
GCCGAGGGATCAGCTGTTAGTCTGATGGGCTTCCCTTTGTGGGTAACCYGACCTTTCTCTCTGGCTGCCCTTAACATTT
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Fig. 6.199

TTTCCTTCATTTCAACTTTAGTGAATCTGAAAATTATGTGTCTTGGAGTTGCTCTTCTCGAGGAGTATCTTTGTGGCAT
TCTACTAGAGGGTAATTTTCTTTTAGGGTGTTATAATTATTTTAAAAATTGATTCATTGGAATATGTTATCAATGTCAA
AACATGAGATAACTTGATGATTATCTCTTTATGAATTTGAGGTGCTGAGTATTTCTTGTTTTTCAATTACAGCATTCTT
TCAAATAGCCAGTCTCATGTTTCTATTTCTCTCTTAGCTGCTTCTTTCTTTAATATTTTGTTGCTAATGAGAACCTCTA
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Fig. 6.200

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Fig. 6.201

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Fig. 6.262

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<p align="center">Fig. 6.203</p>

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Fig. 6.204
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Fig. 6.205

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Fig. 6.206

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GGGCGCCTGTAGTCCCAGCTACTCTGGAGGCTAAGGCATAATTGCTTTAACCTGGGAGGTGCAGCTTGCAGTGAGCAGA
GATTGTGCCACTGCACTCCAGCCTTGGCAACAGAGTGAAACTCTGTCTAAAAAAAAAAAAAAAAAATTTTTTTTTTGCT
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TGAAATGTTGGATGATATCTTTCAGTTTAAAAAATTAGTATTTCTATAAAGTGTCTTTTTTTGGATGGAAATGTTTCTG
GAATGTGACACTTAAATTTTAATTACTATTATTAAATATCAATGGATTTGTTTTGCATAACCAAACAGTTGGATAAGTT
CAGTTTATGTGCTGTGCAATACTAAGATATTTACTTGGCATGGAGAGAATCACTGGCTTACTAAATAAAAGAATTCAGC
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AATAAGATCTGAATTGAGTGTTGAAATGAGTGTTTCCTGAGCTCCAATTAGTGAAAGTATTGAGATTTGAAGAATGGTT
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TTCTATCAAAGGAGATAGATGTACATAAGGGGAATTATGGAGAAGTGAAAATTTGTCTAAGGTTCATTTTTTCAGTAGC
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TCGGGTGAGAGGGTTAGGCCTTTATATCCTTGGTCACTTGATAGTAGATGCTGGAACAAAATGGTGCTCTTTAGCTAAG
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CGCCCGGCCAACAATTATTTTTATAAAATAAAAGTTTATTAATGAATTTATACAAATAGCACTTTATAAAAACTACAAC
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ACCACAAAAACATCAAATTATGGAGAAGTGAGTGCATAGATATATGAAATCAACATTTCAAATTGGTAATAGAATGACC
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TTCGGGAATAGGTCTATCCATCC1TGTCTTTTTGTTTCCTGTCTTGATTCTACAATCTCCCCTTTGGCCATTACCATCT
CAAAATCCTGCTTCCCAAGGTAATCCGTCTAGCTTTTCTAATCTTCTGGCTTCCCATTACTCATTTCAATTCCTTCTTC
TGGCTCCCTGTTGCTTTCCTAACCAATTTCAATTACTTATCCTTATCTTGAAGGCCATTCACCACTCTGCCCTGCTGAC
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Fig. 6.207

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GTGTTTTTCTTATCACCCAAGATTCCTCCCCCGGCTCACACTTCTGCATGCAAGCACGTTCCCAAAGTATGAAAATGCT
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GTGGTTCAAAATTCACCAAGGCAGTAAAATCTTGTGTTTTGTACACAACTTCCTTTCTAATGACCTCTCCCTTTCTTCA
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```

Fig. 6.268

CAGGGTATTTCACCTGAAAGCCTGAAATAAATGTGATGAGAACCTATGAACTAAGTTCCAGTGTAAACAAGTGGGTTAA
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GCATTGATGGATTGATTGTTACTTTTCAATAACTTTTTTCCATATTTGCTCTAGTTTTAAATTTGCAAATTTTAATTCAG
TATTGTTTATAATAAGACAAAAGCTCTTCTTTAAGGTTGGGGCATTAATGTTAAAAAAAAAAAAAAGAACTGTTACATCA

Fig. 6.204

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CCATTGACCATTTCTACTAGGATGCCTTGAAGGCACCTCAGACTCAACAAGTTCAATTTGAAATTCATTATGTACTCCC
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GTAATGGAATTTAGTAAGAATTTTTCTCTTCCTACTTCCCACCAAACAAAAGCAAAATAAAACTAAAATAGAACAAAAA
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ATGTGTGGTGTTGTTTCTGAGGCCTCTGTTCTGTTCCATTGGTCTATATACCTGTTTTGGTACCAGTAGCATGCTGTTG
TGGTTACTGTAGCCTTATAGTATAGTTTGAAGTCAGGTAGCTTGATGCTTCCAGCTTTGTTTTTTGTTTGTTTGTTTGT
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TTCTGTGAAGAAAGTCAATGGTAGCTTGATGGGAATAGCATTGAATCTATAAATTACTTTGGACAGTGTGGCCATTTTC
ACTATATTGATTCTTCCTATCCATGAGCATGGAATGTTTTTCCATTTGTTTGTGTCTTCTCTTATTTCCTTGGGCAGTG
GTTTGTAGTTCAACTTGAAGAGGTCCTTCACATCCCTTGTAAGTTGTATTCGTAGGTATTTCTTCTYTTTGTTGCAAT
TGTGAATGGGAGTTTGCTCATGATTTGGCTCTCTGTTTGTCTATTATTTGTGTATAGGAATGATTGTGATTTTTGTACA
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AATATACAGTCATGTCATCCGCAAAGAGAGATAATTTAACATCCTCTCTTCCTATTTGAATATATTTTATTTCTTTCTC
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GAACTAGCCTTGCATCCCAGGGATGAAGCCGACTTGATCGTGGTGGATAAGCTTTTTAATGCGTTGCTGGATTTGGTTT
ACCAGTATTTTATTGAGGATTTTCACATCGATGTTCATCAAGGATATTTGGCCTGAGATTTTCTTTTTTGTTGTGTCT
CTGCCAGCTTTTTGGTATCAGGATGATGCTGGCCTCATAAAACGAGTTAGGGAGGAGTCCCTCTTTTTCTATTGATTGGA
ATAGTTTCAGAAGGAACGGTACCAGCTCCTCTTTGTACCTCTGGTAGAATTTGGCTGTGAATTCATCTGGTCCTGGGCT
TTTTTTGGTTGGTAAGTTATTAATTACTGCCTCAATTTCAGAATTTGTTATTGGTCTATTCAGGGATTCCACTTCTTTC

Fig. 6.210

TGGTTTAGTCTTAGGAAGGTGTATGTGTCCAGGAATTTATCCATTTCTTCTAGATTTTCTAGTTTATTTGTGTAGAGGT
GTTTATAGTATTCTCTGGTGGTAGTTTGTATTTCTGTGGGATCAATGATGATATCCCCTTTATC`"TTTTTATTTATKG
TGTCTATTTGATTCTTCTCTCTTTTCTTCTTCATTGATCTGGCTAGTGGTCTGTTTTGTTAATCGTTTCAAAAAACCAG
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TGCCTTCTGCTGGCTTTTGAATGTGTTTGCCCTTGCTTCTCTAGTTCTTTTAATTGTGATGTTAGGGTGTCAATTTTAG
ATCTTTCCTGCTTTCTCTTGTGTGCATTTAGTGCTATAAATTTCCCTCTAAACACTGCTTTAGCTGTGTCCCAGAGATT
CTGGTACATTGTATGTTTGTTCTTATTAGTTTCAAAGAACTTATTTATTTCTGCCTTYGTTTCGTTATTCACCCAGTAG
TCATTCAGGAGCAGGTTTTTTGATTTTCCATGTAGTTGTGCTGTTTTGCATTTGCTGAGGAGTGTTTTATTTCCAATTAT
GTGGTCAATTTTAGAATAAGTGCAATGTGGTGCTGAGAGAATGTATATTCTGTTGATTTGTGGTGGAGAGTTCTGTAG
ATGTCTATTAGGTCCACTTGGTCCAGAGCTTAGTTCAAGTCTTGAATATTCTTGTTAATTTTCTGTCTCGTTCATCTGT
CTGATATTGACAGTGGGGTGTTAAAGTCTCCCACTAGTATTGTGTGGAAGTCTAAGTCTCTTTGTAGGTCTCTAGGAAC
TTGCATTATGAATCTGGGTGCTCTTGTATTGGGTGCATATATATTTCAGGTAGTTAGCTCTTCTTGTTGCATTGAACCC
TTTACCAATATGTAATGCCCTTCTTTGTCTTTTTCTRTCTTTTTT`GTTTAAAGTCTGTTTTATCAGAGACTAGGATTA
CAACACCTTTTTTTTTTCTTTCTGTTTGTTTGGTAAATCTTCCTCCATCCCTTTATTTTTGAGCCTGTGTGTGTGTTTG
CACATGAGATGAGTCTACTRAATACAGCACATTGATGGGTCTTGACTCTGTCTAGTTTGCCAGCCTGTGCCCATTTAATT
GGGGCATTTAGCTCATTTACATTTAAGGTTAATATTGTTATGTATGAATTTGATCCTGTCATTATGATGCTAGCTGTTT
ATTTTGCCCATTAGTTGATGTAGTTTCTTCATAGTGTCAGCGGTCTTTACAATTTAATATGTTTTTGCAGTGGCTGGTA
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CCATCAGGTTATTTATGTTCTTCTCTAAACTGGTTATTCTAGTTAGCTATTCCACTAACCTTTTATCAATGTTCTTAGC
TTCCTTGCATTAGGTTAGAACATACTTTTTTAGCTTGGAGGAGTTTGTTATTACCCACCTTCTGAAGCCTATTTCTGTG
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TCTGTTTTTTGGAATTTCCATCCTTTTTGCACTGGTTTTTCCTCATCTTCATGGATTTATCTACCTTTGTTCTTTGCTG
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AGCCAGAACTCTCCTGTATGAAGTGTCTGTCAACCCCTGCTGGGAGTATCTCCCYATCGGGAGGCACAGGGGTCAGGGG
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TTCCCAGGTGGGGAGACGCCCCACCCTGCTTCTGCTTGCCCTCTGTGGGCTGCACCCACTGTCTAACCAGGTCCAGTGA
GATGAGCTGGGTACCTCAGTTGGAAATGCTGAAATCACCTGCCTTCTGTGTTGATCTCACTGGGAGCTGCAGGCCGGAG
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CAAGCAGATGGCATTGCTTGATTAGAGTCAGGCCCATATTTAGAACATTTCTATAAAGCCATTCTCATTTGGGCAAACAT
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Fig. 6i2I1

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GAAGCCTCTTTGATGAAGGTGAGGGTCTATAAACATGCTTGTATTCATTATAAGGGGATTGTGATTGTCCTCATAGAAA
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CTGGCCAATTTTTTGTATTTTTGTTTTTTTAGTAGAGACGGGGTTTCACCGTGTTAGCTAGGATGGTCTCAATCTCCTG
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AATGCCATAAAGAATCCATGAATAATATAGTCGTGTATTTATTTTTGAAAGAGAATGTTCTTTTAATAAAGCATATGTT
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Fig. 6.212

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TTCTAATTTACACTTGAGCAAACCTTTATTTTAAAGAAAAATTGAGAAATTTATGTTTCAGAGTTATGAGACACCGTCT
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GTAATAGGAGTCATCTTAGAATTGTGTATTAGCCAAGATGTGGAATTCCTATCAATAAATTAAGCAGTGTTCCATTA
TATTTGAAAAGGAGAGAAGATTAGGGTGGCTGATCATCTTGTATTTAAATGCTAAGAAAGACAATAATCACAGCCAARA
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CGGGGCTGCTTTATATATATTTTCTCTTCTGTAAAGTGATAAAAAATAGTAAGGCCAAATAATATTTATGATAGTTAAC
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AATCACTTGTTAAAAAATAGCATTGCAGAGAAACATTGAACCCTCCTTAAATGTAAATAGCACGATTCATYGAACATCTA
GTCAAGATGACTAATCTTTTTTTTCTAGTAACGCTCAACAGTCTAGAAGGTACAGAAGGCGGTGGGTATCAAGGTTTATA
AAAGGTCAGGAATTGGAACTCTGGGGACCTGGTTGTTCTCATGAGGGCACATACCAAGATCAGGGTGAGCTAGGAAGTT
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TACATTCGTATATATAGTGTGTGGGTATGTGTTTGTGTGCATATGTGTGTGCATATATATATATAAAGAAAATAGTGTA
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TAAGGGCAGGCCCTAGTCTAAGAAACTAAATCTAAGACAGCCACCTGAAGTGTGAAGATGGAAATTGCAGAGTTGAGTC
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TTATTTATACTCCTACTCTAGGGGGCCACAAAAAAAAAAGAGGGAAATGAGAGTATAGGAGATTTCTCTAGGTTATCATA
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Fig. 6.212

AGAAAGAAGTCTGACAATACAATCCTTGTTCCATTTATCTTTTTATAATAAACCACAAAATATTTAAGAGCTGTTGGAA
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AAAATATCCCTACAATTCATGACCTTGAGTGACTTCAAAATTCAGTTAGACAAAACAGACAGCTGAGTTCTGTGGTTCT
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GCTCCAACCAACCTCAAAATCTTAAACTCAAATATCCCACACTCTAAGCATAGTTTTCTTTCTTTCTAGCTCCTCTTAT
TACGTTAAAGACCTAATAGTCTTATTTCTCAACTGCTTTCTTGCTAACCCTTCATTGTTCTCCTCATAAACCCAAACAG
GAGATAACAAGCCAGGATTAATCCACCTGTGTGATCTCCCCATTTTTAACCCTTGTGTTTCCAGAGAAAAACACAACCA
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CCTTTTCTCCTCCCCATATTTCAGTCTGAATGAGTGACATCAGATTTTATTAAATTGTACAAGCCAGAAACTTGATAA
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GGGCTTTCTTGTTGTGACATCATATGGCGGAAAGTGAGAGGGCAGGAGAGAGCAAGAAGGGACTGAACTTGCCCTTATA
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Fig. 6.2[4]

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Fig. 6.215

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Fig. 6.216

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GAGTCATGTTCAAAAAATCATTGCCTAAATCAGTGTCATATAGCTTTTCCCCTATTCTTACATTTATGTCTTTCATCCA
CTATAGGTTGATTTTTATATATGGTATGAGATAAAGGTTTATATTTATTCTTCTGCATGTGGATATACAGTTTTCCCAG
CACCATTTATTGAAGAGATTGTGTTTCCCCAATGTATGTTCTTGGCACCTTTGTTGAAAATAAGTTCACTGTAGATGTA
GGGGTTTATTTCTGGCTCTCTATTATGTTTCATTGGTCTATATGTCTGTTTTTATGTAAGTACGATGCTGTTTTGGTTA
CTACAACTTTGTTGTGTATTTTGAAGTCAGGTAATGTGATGCCTTTGGCATTGTTCTTTTTGCTCAAGGTTGAGTTGGA
TATTTAATGTCTTTTGTGGTTCCATATGAAATTTAGTATTGTTTTTTTCTGTTTCTGTGAAGAATGGTATTGGAATTTT
GATAGGGCTTATAATGAATCTGTAGATTGTTTTGTAGATTGTTTCAGATATGGACATTTTAATATTAATTCCTCCAATC
TATGAACACGAACTATCTTTGCATTCATTTGTGTCATCTTTAATGTTATACAGCTTTGTAACTATTATAAATGGGATTT
TAAAAATTTCTTTTTCAGATAGTTCGCTGCTAGTGTATATCAACACTACTGATTTTTGTATGTTGATTTTGTATACTGC
AATTTCTATTATAATGAATAGAAGTGGCAAGAGTGGCCATCTTTTCTTGCTCTGGATCTTAAAGGAAAAGCTTTCAACT
TTTCCTTGTTAAGTATGGTGTTAACTATGTGTTTGTCATATATGGCCTTTGCTTGTTGAAGTACATTCCTTCTATACTT
AATTTGTTGAGGGGTTTTTATTATGAAAGGATGTTCAATTTTGTCAAGTGCTTTTTCTGCGTCTATTGAAATGATGATA
TAGTTTTCGTCCTTTGTTCTGTTAATGTGATATATTACATGTATTGATTTGTGTGTGTTGAGCCATCCTTGCATCCCTA
GGATGAATCCCACTTGATCATGGTGAACAGTCTTTTTATTGTGTTTTTAAATTCAGTTGGCTAGTTTGTTGAGGATTTT

Fig. 6.217

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TGCATCTATATTCATCAGAAACTTTGGCCTGTAGTTTTCTTTTTTTGTAGTGTCCTTCTCTGGCTTTGGTATCAGGGCA
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GGAGAAAGAGCTGAAATTGGGGTTCAGATTTATAAATCTCAAGTTTTGGTTGAGCACGAAAGGAGAGGGAGTGAGACTA
ATGATAAGAAGGTTAAAGTAATCACTGGATTGATTTGGGATGGGCAGTCTCAAAAGTGCTGAGGATTCAGTTCCTATTT
CAGTGAGCTGGAACACACAAACATAGAGGAAAAAGAAAGCAGTTCAAGTATAAATTTGAATATACCTAAATGTGTGTA
AATTACGGAAGAAACAACTTCAAAAGAAAATAACAACAATGACATGGGCTACAGAACAGTTAAGATTAATGATCCCTCA
TTATAGGGTGCCATTCAGACATAGGAGTTAAAAAAAAAAAAAGCAAACCAGAACCACACACTGCAGCAGCTCCCCAGTT
GCAAGCACACTGAAATAACAGAGGCCTGCCTTCACGGGAAGAGGATTAGTTCTACTGTGACCTGTTTATATAAAAACAA
TGTTGTGGGTAGCAAGATTACTAGAAATGACTTCTTGGATTATAAAGGACAATTATGATAGCTTAGGCCACTAGCTTTC
AGTACATTCCATCTGGTACGTTACAAATTAGCTCATTATTCTGCTGTTCTTACGAATGTGTTAGTCTGCAAACATATCA
TTGTCACAAATTAGTCACCTAGTGTTTCTCTGAAAGTGATTTAATATAAGCACACCCACATCATAGGATGGAACAGTAA
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TGCTTGAGTCAAAGGAGTTGCAGCCAGAGGACCCAGCAGGGATACCAATTATGAACCCACAAAAGTACCATGAGACAAA
GAGGCAGCTTTCAACATCTGCCAGGTCTCCAACATGTGATGCCATCTTACGATGGCACTAATTCAGGAAAACAAAACAA
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AATGGGCAACTAGTATTTAATAACTAGCCTTGAATTTATTCTACACTTGAATGCAAAGATTATTCTACTTAATTATACT
ACAGTGACTTTGAAACTATTAATTATTCTAAATTATTGGGTTCTAGTTTAAATACATTTTTATATTTTTAAGCTTTCAA
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CTCTATAAATCTAAAAATATCACAGTATTTCTTTTTTTTTCCAAAATCTCTCTCTCTCTCTTTCTCTCTCTCTCACCCT
CTCCTTATCTGAAACAAACTGACATTTAATGAAGAAACTGTGAAAAGGGGTTCGAGAAAATGAGAAGCTAGTATTTTG
ATTGGAAGTGTAATAGATGTCAGTCAGAGAAAAGAACAGTCTCCAGAACATCTAGATGTGGATGTAGATGTAGATACA
TTGTTAACGCCACTCTGTGCATTTTAATTTAAGCCACTTGATCATGTAACCATTAATTCAATGTCAAGTTCATTCATTG
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Fig. 6.218
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TAGCTATTCAATATTACAATATAGCATTAATTTTAAATGACTTTATTTCAGGAGGGAAGGAAATAATTTATGAATAGCA
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ATGGTGAAGTGTAGGAAAGGCAAGGTCTTTAGAAGGAAACATAGGCAGACATAAGTAATCATGTTTGGTATAGAGACTC
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ACCAGATACTATTCTAGGCATTAGAGGATATACCAAGGACAAAACAGACCAGAAAACCCCACTACCCCTAGAGCATGTA
CCCCACTACATGCTGATGATAATCTTTAGTCATCTATTTCAAAAATAACTTAAAAATGGAGTGGCAATGTCTCTGCGTA
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GTTGAACTTGCATGGTTTTCTAAATGAAATATTTTCATTTTTAATGTTTCATATATGTGTCTGTATATATATATGTAAT
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TATTTTATTTTTGTATTTCCCATGTTGAACTTGCATGGTTTCCTAAATGAAATAATGTTTCCTCTTTAAAAAGAAACTT
TAAATGATATTGAATCTAGTTAAAATGAAACTATGTGATAAGTTTCAGATTTTATAAATAGATTGTCAAAATATGTGTT
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AGCACTGGTTCTCTTGGAGGTTTCTGCTCCGGTATGTTGTGATTCTCCATAGCCTACTGTCTATCTCACCAGTGGTTTG
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Fig. 6. 219

GTTGAATCATGAAAAGTCCCAACAAAGCATTTGCTGGTTTTAGGTGATGAAACTAGGTGTTTAATGAATTTATTTTTTC
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CCATAAAGATAATCTTTTATGGTTACTCCTGATATTTCTTTTTAATTTTTTTAGAGACAGAGTCTATGTTTCCCAGCCT
GGTCTTGAAATGGGCTCAAGTTATCCTCCTGCATCAGCCTTCCAAGTAGCTAGGACTACAGGCATGAGCAACCAGAACT
GGCTTCCTGATATTTCTTAAAAGTTTCATGAAAGGATGATTATCTTTTAGTGTAACAAAATAGATTAGCATTACCCTAA
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GTAAACAAGGCTTCCCCACCCCGGTGCTTTTTTTTAAGGAGTGAAATCCACCAAACTCTATCATTTGCAAATTATCTC
TGGACTTCTTTTATTTCTTTTTTTTTCACTGGGTAAGAAGGAAAACGAGCAAACAAAAAATCCTCCCTCCGTATTTGTT
TAAGGCGTAATTTCCCTGTGAGGTGTTTTACAGGCTTCTCCTTGTTTAGCTTTTCTTGCCTGGTAATTACTGAAAGAAG

Fig. 6.226

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TCTGCTCTTCCAGGGCGAGCTATTCCTTTAACCCCTTTTGCTTGCTCTTACTTGGAGGTTAGCAAGGAGTGCGCGGCTC
TGCAATGTCTGGGAACCCACCTAGAGGGCCCCCTTGACGCCAGCACATCCCCTGTCAGGCAGCAGTCTCTGCAGCTGAG
CCAGGCTGCTGCAGAGTTAATGTACAGTACCACGGAGCCTGCAAGTGTCCTGAGCTGATCAGAGCTGGGGCGGCACAGC
CCAGGGCAGACAAGGCGGCTGCGAGGATTCCAAAGGTTCTGCTGGAAATTCGGCGCTGGGGGACTCCAGCAGGAGCCTG
TTGCCATTGTGTTTTAAAAGCATGCAAGGTCTGTATAGCTTGGGCATGAGAATCTTTCAGAAGATGTCAGAGCATCAGA
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CATCGGGGGGAAGAATTCTTAGACAACGTCTGACTTGCAGCTGTGAAATTCATTTTCTTTCCAAGGCAAAATATGAAAAG
CTCCCACATGGTTTTGATAATAACAAAATAAAGGGGATTCTGCAGTGAAAAGATCAATAGCTTAGTCATTTTACTTAAA
GAGAACAGCCAGCCTTATTATGGGGTTAGGCAGCAGAAATGAATTTCATCGTGACAGCATCTTCTGAAGTCATGATGGT
AGTTAATGGTAATCTTGTCCTGCAGAGCAGAAATTACTCATTGCCTTCCTACTTTTGCAGTTGAACCAGCAATTCTGAA
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GTAGCACTGCGTTAAAGTGACCATACATGAATATGTGCCTAAGAAAGAAGAAAAAAGCAATATTCATTTCTAAAGGACAG
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TGGGGGGAGATTTTTTTTTTTTTTTTGGGATTCAATACTTGTTGCAATAATTGCCCACGATAGCTGCTCAAACAAGA
GAGTTGGAATTCATCTGTAAAAATCACTACATGTAACGTAGGAGACAAGAAAAATATTAATGACAGAAGATCTGCGAAC
ATGATGCACGTGAATAATTTTCCCTTTAGAAGGCATTCCTGGATATGGTGAGTAATCAATATTCCCTTCAGTTTGTAAA
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TCCCCAAATAAGTCAAGCATGAGGACTTGAGAACATTTAAATTGCTAATATTTCATGGAGGAAGAAAAAAACTTTGAGA
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TGCTTAGTGCCTTGATATTTGGTCATCTATTAAGGTTCAATGCAATGCATTCAGGTCCCTGGACATGTGATATCATGTG
CCATTTCTTGCTGGAGTTTATGACTAAATGTGTGTAGGAACTCATTGCTGAGGGTATTTATAAGAGCAAAAGGTCATTA
TTTTAAAATTGCTTGTTTTTGCCATACCTTTAGGGTCAGCTGGATTCTGACTTTCATAGCAGAAACTTTGTGAATGCAT
AATAAAGGCACATGTTAAGGCTTAGTGTCTGCTAACATGGGTTGTTTTGGAAATGCAGTTTGTCTGATTTTGAAAGTAT
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GCTTTTATGACAATGTCTCAAATGAAAGCTCCCTGAGAGCTTAAGGTACCACCAAAATCACCTGCTGGTTTGTTACAGA
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TCTAACAACACCAGATCCATCCAGAGTCGATGCTGTGGCGTATCTACCTTTTTTGCTGACCAGAGCTACTATCCCCAGT
CTCTAGAATGCTTGGGTGACATGCCTGCAAACCTCGGTGGCCCACTTCCAACTGCATCACCAGAGTTTCCTAGTCAGGG
GGAGCCTTGGTGCCATTGCCTTGCTTGTTCTGTTGGTGAGGGTCAGGCATCAGCAATAAGGTCCTCATTATTCTTACAG
ACAAATTTACATCAATAGTCTTTAATCTTGAGATTAAAAGATCCTGGAAACAGTTCCTGGCACGTAGTAGGCATTTGTC
AATTATTTTTCTTCTATGCCTTAGGCTTTTCTTCAGAGTTCATTTATACCTCTTAAGATTTGCTTGGGAGGGGAAATT
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TAGATAAGTACTATATAGCTCTTCTTATTTTTCCTTCTGGCTAGTTGTTAGAATGGAGAGATAGCCTGGCATTCAGGAA
CAAGTATGGCATGGTTGAAAGAAGGGAAATGCAAGTCAGCTTTCTAGGAATTTAAATTTCATGTAGCAGCAGTTAAGAG
GACCTTTAGAAGCATTATGACCTGGAATCACATGCCAGGGTCTAACATGAATGACCTAACACAGTGTGACTAACATGTT
CAGCTTTTTGCCTGACTTAAAAGATATATAACAACTTTTTATAATCTTATTTGTAAAAATACTATAATTTCATGCCAGA
AACATCAAGGCTATGTTGAATGCAATTGAATTTCATAGCTAAACAATTTAGGAGGGCATTTTATTTAAATACCTTTGCA
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Fig. 6.221

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TCCTTTTAATGATGGATTCACATGGCTAAAAGGACAGTGGAAGATAGAAGCTAAAGGCCCTGGCTAGGAGAGGCTGTGG
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GCTTATTTGGTGCTTAGGGTTTAATTTTTTAAAGGTAGGACATCTAAATGTTTTAAACTGTCTTTGTGATGCTGAGAGGT
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CTCATAGAGTTTCATCTTTTAAGGTTTTCTCCTCTTCATACATTGAAATTACAGAAACTCAATCCAGAAGTGCATTTTC
CAGAGTATGTGTTCTGACCAATACATTATATTTTATTAGAATTTTAGGGGGTGGGAGCAATATTGAGTTTGATGACTAT
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TGCAGTGTCTGCTTGTCTTCCAGTGAAATTGCAGGTAATAGCTGACCTTGCTTCTTTAACATCTCTTCCTCTGGGTTAC
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TACTTGTAAACACTTAGGACCAATAAAGTTTAGATGGAGCTCATAATTATACAAACTCATCTCGTTCACAAATCCCTAG
GGCTCAATGTTAAAGTCAGCCATTGTTTAAGGCAGAAATTCAGGTTTAGATATAGTGTAGCAAAGATTTTCCATTATAT
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CCCTTAATTTTTTCTAGTAATACCTTGGAATTTAAAAATGAAATTACAAATGTTTATGTTTTAGTGTTTTTAAAAATAA
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CAACTAGGGATTATGTTGTGTAAAATTTATACATATATATCTCTCCATATATGATTTCAAGGATAGATAATCTAGAGTC
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GGGAGAATGCAGATAATGTGGAGATTAGGTAGGTGGAAGGAGAGCCTGAGGGAGTTTTTGTTTTATGCCCTGTGTTTTC
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ATᵀᵀᵃTGGAACAGCTACTAGGGGGAATTGGAAAGGTTATTGAAGAGAGAGTACAAGACAGTCAAACAAGTTGGAAAAAGG
TᶜᵗᵗᶜᵗGATGTTGAATGAATATATCCAGTCAATAAAAGACGTTTTACTCTACTTGAGAAATTATTTACACAAGTATTAT
GᵗᵗᵗᵗGGAAATGAGTAGAGATGTGACAGAAATGTTACAATTCTTAATTATTAAATTTGAAACTGTTTTGTGATGAGAAT
GTᵗᵗCCTATGGATTGTGTTTCTTGGGATTCCAAAGAATACAGAGTTGATCTTAATTGTTGTTGTTGTTTTATTTGGTAG
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Fig. 6.227

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TGGTGGAAACTGTTTTATGAACATATTAATTAGATTGCAGTTCGATTTTCATGACCTGATTTTTTGAAAATGTGAGATT
TCCTGTGCTGCCTTGAAATAACTCTTGGGTGACAAACAGAAGCTTTAAGATGTTTGCTTTGAAT. . TATCTTCAAATGA
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GGGAACTTCATTCTTCCTCTCCCTTGAGTCTCCATATTTTTCTTTTCTAAATATGAAAATGAGAAGGTTGACTTAACTG
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CCCAAAGTGCTAGGATTACAGGAGTGAGTCACTGCAGCTGGCCTAGAAGAGCTATTTTGAACGTGCCCTTGGGATAGGA
AGAAAGTCCCTGAGGATCTGACTATAGGTAGGTGCTCAAGTGCTCTAAGGGTGAAAACTTCTCTGCATAACAGAAGAAG
AATTTAGTGCTAAAGAGCCAATGAGAATGATGGAGATGAGGCAGAGAAATTAGGAGAGAGGACAGGATGAGGAAAATTC
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Fig. 6.223

GCCCTGGGAAGGCTGAGAACTGCGGGTATACTCCATAAACAGATCCCTCCCCAGTACTGACAGTGGTTCGGTGGGTGAG
GGGTATTGGGGAGGGTACAGGGAGTTGCTAGGGGTGGGAAGACCACTAATATATGTATGTCTTACTTGAAGCCTCAGGA
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CAAATCAATTTTGTTTAGATCTCAAGGGAGGCAGAGGGCCTTCCATCTTTCAAGAGTTAAGCAGACCATGAAGTCCAAG
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CGTGTCCATTGGATAAACAATTTGATGGATCAGGGAAGCACACCACTTATGATACTAAGATCAGAAAGAAATTTGTCCT
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GCAGTAAAAGAGATTAAGCCCCAGAATGAGTCTGGATGTTCTAATGGTCAGAACAGGCATTTCATATGACATGGCCTGC
ATTGGTCCAGGTGGCGAAGTAAGGATGGACAAACATGCGTTGTTTGCAATGATAAAAAATCATAAGTAGATGATAACAT
TGATTTGCTGAAATTTTTCTCTTTTTTGTTTGGTGTAAAACTTTATATGTCGTACTTTTTGCCTTGTGGTTTTTTTGTA
TCTTAGGACAAAGGTATTCTTAATCCCTTTCTTTGTCTTCTTGAAAGATGCAACCATGCAGAAATGCATACCTCCTGT
AAATTGTTTCCTGTCTTTTAGTAAACCTTTGTTTAAATGAGTCATATGTTTAGCAATTTTAACATTTTAACTTGGCTCT
AATTAATGATAATATAGCGTGATTTATGGTCTTCTAAAGAAATGTAGACATAGAAAATGGAGATAGAGGTTAACATGTG
AAGAAGAGTTTGTATTTTGGGTTAGAAAATTAAATATTGGCTTATGTCAGTAAGATTAGGCCTTACTATTTGATTGTAC
GAGTTTACTTTAAATTTCTAGGCTTTCAGGGCTTTGTAAAGATAAATTTTATTTAAAACAGCTTTAGATTTGCAGAAA
AATGGCAAAGATAGTGCAGGGAGTTTCCATATACCCCTCACCCACTGCCTTTTATTAAAGTCTTATATGAGTATGATAC
TTTTGTTACAATGAATGAACCAGTATTGGTACATTATTATTATTTAAAGTCCCTATTCATTCATATATCCTTAGTTTTA
CCTAAGGTTCTTTTTTTCTATTCCAGGGCCCCACTCAGGATACCACATTGCATTTAGTTCTCGTGTTTCATTTATCTTT

Fig. 6.224

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GCTATAGAATTTTCTCAGATTTTCCTTGTTTTTCATTACCTTGGCAGTTTTGAGGAGTACTGATTAGGTATTTTGTAGA
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CTCCAAGGAGCACACGTTCCTTTCATTGGAAAATGGTATTAAAAACCAAGATTTGAGCAATAGGTGTGTTCATTGTTTT
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CATATCTAACAATCAGGAAGCATAAATAGATAGATGATAGATAGATAAGCATTAGTTCATACTGATATTTTCAAATCTA
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TAAAATGTATTTGTGTCCCAAATGTTTTTTCTGTACATAAAAATTGATATAGTCAACTTTATATTGTTTGCAGTTAACTT
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CAAAGAACTACATGTTGACTGTATCTTGCTGGTAACTTAAGGAGAATAGTAACTTGTAGTTTCCACCTTTAGCAACTGG
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CAACTTTTTTTGACGTGGCTAGATGGTTAGGAACACACTGAAGAATAGATTTTCAGTAATGAGCCATACTTTCAATGAA
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CCCCCCATAGCCAAACATTATTTGCATTCATCACTCATTCGTAGATGCCAAATCAGACATGGTTCTTGCCCCTATGGAG
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CCTTTCTATCTCTCCCTAGTTGGATAATAGTACACATATTAAATGTGTACCTTGTACCAAGTATGCTGAGACCTGCTTA
TAACTAATTTTGCTTGATCTTCATGAATAGTGTTTTGAAATAGCATTTTTCTATCCCTGCAGATAGGAAAACTAGAAAA
CAAAGAATAAGTAACTTCCCCAAGCTCACATGACTTGTAAGTGGCAGAGCTGGGATTTGAACCCAAGCCCCTCTGAAGC
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Fig. 6.225

TAAAGCCTGTAGAGTACATGCTCCCTTCCTCCACATCACACTGCCCCTGCTTGTTCACACTCAATGTGAGAATCCACTT
CTCAAGTTTGTCTTGCTTATTCCAACCTGTTTTGCAATGAATCTCTTTCAACAGCTCTATAGTACCTGGCACTCACTTG
GCATCCATCATTTTCTATGTTGTCATGTTTGGTATCTTTTCATGTATGTAATGTGCATCTCTTCTTAGGAGTAAGACCA
GAACTTTTCTGCTCTGCAGTTTTCTCCCTGTGAGGGATGGTGAAACTTGGTGGCTAAGAGCATGGTTTATACAGACAGA
CTTCTTAGGTTTATATTCTGGCTCTCCTCTGTACTTGCTGTATCAGCTAGGGCTGATGTCTTCACTTTTTTATGATTCC
ATTTCTACCTCTGTACAATAAGGCAAATATGAATACTTACCTCAGAGTGTTGTTGTGAAGATGGAATGAATTAATGGCC
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GCAGATAATTGCATTCTTATTTGAGAGTTATGTTAACTAGTGTGGGCTTTTATTTGCTTCTCTCTTTCTATCAATGAGT
CCTGCTGCTTCAACCCTGGGTTAACTGAAATTTGAAACCTTTTTCTACTCAAATTCCTTTGATGCTGAAATTCCTCTGC
GGTCACAGCTAAGCACTTTGTGTGGTAACCTTCTCTTCTGTCCTCAGGCCTGCTGAGATCATCTGCTCAGCAGCATGTC
TGACACCTGTTGCCACCTCACCCAGCCACTTCCTTGGTGACCTGGCAATGCCCAGAGGGGCCCTGCCAGCTTGGAGCCC
TGTGGCAATGCTGTTTGATTCAGCAAACTGATAGCCTTCTGACATTCCTTTCTTCATTTTAGCAGCCAAATCACATGCAC
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CAAATAGCTGGGATTACAGGCATATGCCACCATGCCTGGCTAATTTTTTTGAATTTTTAGTAGAGACGGGTTTTTGCCA
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TCCAAGAGGGCCACTGGAGGCTTAGGGGCTTTGAAGTTAACAGCATGAAACTTCTCCAAAACAAACATTTAATTTTTTT
CTGTTCGTTTCCTTAAACAAAGTCCATGTAATAAATGAAATAATTTTGAAGAACATTGTAATGAATCCCACATACCAGT
GGGTCCCAAATTACTGGAAATCATAACATGGCTCATGTGAAGAAAGAATTATACTATTTTCAAGCTGTTGTATGTTAAA

Fig. 6.226

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CTATTATAGTAAGTAGCATAATTGGCATATATTTTAGGCCGTATTTTCAATAAATCAATATTTTTGCTCAAAAGGCAAA
ATTTAATAATTTAATAATCTCATAAGGTGAAGGATGCACAGGTACCATGTTAGGAATTTGACATGTGTTGTTTGCTAAT
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CACCCTCTGGTAGGCCCCAGTATGCTGATCCTCTTTGTGTCCATGTATTTTCATTATTTAGCTCCCACTTACAAGTGAG
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CCAGTCGTTTCATGGTTAACTTTCCATTCAGTTTCAGTTGTATTTCATTGTGTTTCCTGCCCAGTGGAGATTCAGCATA
GATGACACCTAAGGAAAAATGCTAGTCATAGTGTGAGAAATCATAGCCTTCTCTTTCAAATGCTTTTATTCATATTCCT
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Fig. 6.222

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TTGGACTTGATTTTAGATATTGCTCGCCAAATTTAGCCATCTTTCTTGGTAGACATTTCTGAAAGTTTAGTGATTAAAA
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CCTGGCATATGGGAGCTATAAACCACATCAGCCCTCCTTTGCTCTTTTTCCTTAGGGCTGCCCTGAGGAAGAAAGCTAA
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CTTTAGTTCTTCCACATTTCACTTTGTAATGTTTGCCTTGCCTTGTATTTATTAAACAATAGTTTACTATACTCGTTCC
TAAAATATTAAGTATAGCTTATTATGCCCTGTTAATCTTTCTCAGTGGCATACAATAATTTAAATATCTGCTTGAAAAA
TCTCACCAAGCATTCATTTTAATGTGGCTTAGAATGTTATATAATATGAAAAAGTACGTGTTCTTTCAGAAATTAGTTC
CAAAGAACTCCCTATATCTAGAGACAGGCTGTGAGAATGGAGAAAGTTGAGGAGTGTCTCTGTTAGGAATCTTACTACT
GCTTTTAAATTTCATGGAACACCTATTAAATTGCTTCGCAGCTAGGTTATGAGGAAGGGGAGCCCTGCTCACTCTTTTA
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GCCTAGGATGAGTCAGAAACATGTACATACGGAAAGGGAGAGAGACAATCCTGATTAAATTGTGCCTTCTTGCAAAAAT
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CAAATAGTATGTTATATAAATTTTCTAGAGTTTGTGAAAAGAAAATCAACTTTTCAATTCATTCTATCAAGTAGACCTA
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CTTGCATTTTGTTAAAAGATTCATGCTCATGTCAAATAAGGCCGAGTCTGTAAATGTACATATAGTTAATTTTGATCAA
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TTGACTAAAACTTAATATCCATTTGTTGTTAGTTTTTTTTAAAAGGATGCAGTGAAATAGCAACAGAAGAATATAGTGT
TTATTTCAATGGTGAAACCAATTTTTAAGTGTGAGTCGTTTCTATTAAAGTCAGCTTAGTTATAAGCAATAAGACGTAA
CATAATATTTGTATGTAATATTTGTTTGGGAAGTTGTATAGTTATTGTCAAGTTATTGTTCTAGTACAACAACTTGAAA
CATAAAAATATAAATGAAAACTGACTTTATAGCTATGTGGGTATTTACCTGGAAGAGCCAAAGGAATTAGTTGAAATGT
TATTAGAAATGATTCAGTAATTCAGATAGGTAGCCAAATACAAAATTACATATAAGAAAGCTATTAGCTTTTTTATAGA
AAGTAATAATCAGTTAGATATTATAATGAAAAAAATAAAATACCTCTTTATTTACTTACTTTCTTACTGTTGTCCAGGC
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GTTCCCTAGGCTGGTCTCAAACTCCTGGGCTCAAGCAATCCGCCTGTGTTGACCTCTCAAGGTGCTGGGATTATAGGTG
TGAACCACTGTGCCCGGCCAATTCTTTTTAAATAGCACAAGATTCTAGTCACAAAAACATACTACACCTATATTTATGA
GTGAAAATAATAAGAAATTAGAAGGGTTTTTTTGGTATTGTTGTGAAGAAAACAAAAATGTCTTAACTAATGCTAAACA
ATCCACAATATTCATTCCACTATAGGCAATTCCAATCACATTTTATTTTTGATTTTAATAAAGTATTTCAAAAATCAAG
CTAACATGGGAAAATCAAGAAAAAGTTGAAATAAAGCACAGTGGGTAAAGGTTGGAAGGATGTAGAACACACCATTCCA
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GTTTGGGAAAATAATATAAAGTTAGATTCTTTCTTTAAACCTTATATAAAAATATACATTGGATATATATTTTATTAAA
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ACGTAAAGGCAAATGAAAAGGGTGGGATTCTATTTCCAGTGTTACTAAAGAGACATTGACTTTAATGTCTTAAAAATCC
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ATGCATAAGGATAATTTTTAACATGGTAGATTTTCATTATGTTAAATGAAAAAATAAAAAATACATAAAATGAGAGCCC
ATTTTTATAATAAAAAATATATAGTCATTCAAGGAAAACATGTAGACGGACATCACAAAATACCAACAACAGGTATGGT
TCAAAGTGTTATAGGTGGTTTTCAGCACATAGATTACCCCTAGCAAAGAGACCTCTGTGACACTTCGTTTCTTCATCTG
TGAAATAAAGGAATCATACTAGATTAGTTCTAAATGCTTTTAGCAAAAATATTTCATAATTCTATGAGTATTGATGGTT
AATAACCTTGAGAAAATTGTTATTATTTATTGAACTTTGGAGTTAGTAGAGTAAAACCAAGTTTATCCTGAGCTATCCC
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Fig. 6.22b

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TTGTTTTAGTTAAAATTAGTTAAATTTAACATCCTGATGTTTATTTATAAATTGTGCATGTGTACATGTGTGTGTTTGT
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GAACACTTAACATTTCACAGTGGCTGGCTGTTTGGAGGCCATATACAATGGTAGAAAAAGTAGCTTTCTAAAAGTGCTT
TGCCTCATTCTTGCTTAACCTGGCAGACGTCATGAACTTCCAAGAGGGAACATATTTTATTTGCCTTATACCACCAGAG
GACTTGGCCAATTATTGCTATTTATTTAATGTTGGTGTTGACATAGTGCCTTTGCCCTGAGGAGCTCCAGGTGCTTTAT
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AAAGTTCTAGTGGGGTGTCATGGGAGTTTCTCTCTTGTGTAACTAGGTGAACTCCTGTAAGGAAAATTTGCCATGCACC
CTGCAGAGCTGAGACTTACTATTCTCTGTTGTAGGACCCATACTTTCCCCAGTAAGGAACTCCATTTTTTTTCAGGCCT
TTGCTCAAATATCACTTTCTTAATAAGACCTTTCATGACCATCTGTTTTTAAAAACAACTAATCAAAACCTATCCTCAA
GTGCAGACTTTCTGTATCCTCATTCCTGCATTATTTTTCTCTGTAGCACATGTAGCTATTCAGCACACTTAGATTTTTA
TTTATTTATTAATTAAGGGTTGACTCCATCCCACCAGATCGTAAGCTTTATGAAGGTAGAACATTTGTCTATTCTGTTG
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TCTATGTGGTAGCAAAGAAAGCTCAGACAGCACAAGGCTGCCTGTAGTAATTCTTGTGACATGAGAGCATACATCTCCC
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AATTCTTTATCAGAATAAGAGGAGAAGATGCATGACAAGTATTCAGAGATACACTGTAACCAGAAAAAATGGTCATATG
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TGTCATAGCTAAATTGGTTACATTTTAAGCTGTGAATGAAAGGGAATTTGGTAAACCAAACAACTGTTGTATGCTTATT
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CAGGTTCCAGGCACTGTTCTAGGCACTGAAGATGAAAGATGAATAATATGGAGTTCCCATCTTCAAAGAAGTTACAAT
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Fig. 6.229

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TGGCTGGTCTCTTTACCTCTGAGGTTCTTTTGCTTTTTATGTAATTGGATTTTAACCTCAGAAGACTGTAAGGTTAAAG
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TGTAGGAATTTATTTGTTGCTCTGAATGTTAACTAAAAGTAATGGGCTTCTTATTTTGTCCTTTTCCTTTTGTGTTATA
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ATTCTGGGCACGTTGTCCCCTGAGTCCTCTGAAATCCATGTCAGAATAGAAGTGCTCATTACCAGACTCGATGAGGAGC
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CAGTCTTGGAACTTTTGCCCTCTGGCATGGGTAGAACTAGCGTCAATTGGGTAATTCAATCCCTGAGTTCCTTCTTCAA
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GCTCAAAAAATTCAGATAGTGTAATCTACTTTCTCTCTCCCTCCTCCAGATTTAAGTGGATTAATTCACTTCACTATGC
CAGCACCTGAGTACAGAATAATTTATTTCTGTGTCGTTTCATGATGAAAGTCCAGCTGGCATTAGCTCAAATTATGCAG
CTGGCTATTAATTGAAGAAAACTTCCCTGTCACTCCTCAGTCAAACATGATCTGATTCAGCAGATCCTTTTCACTCTCA
TCTCTTCTATTCTACCAGTAGAATAAATGTAAATAACTTTCACTTGGCTCTGCTTGTCATGGATTGTAAATTACCTGGC
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TGGATTTTTAACTTGCTCTTCTCAGCACAATGACACAACAGAGGACACAGAGGTTAATGGAATGTCTTTGGAGCAAAGG
AGATTTCAAAACCACCAGTTTGTTAGATTTGATGATCAGCTCTGCAAAATGTTTAAAGAAGATTTCTTACTCATATCTT
ATAAGTTAAGACTATAAATGGTTTAAATGGGGGAGGATGTGACAGGATTGAAAATCTTCCCCTTCCTTTGTGGCTATGA
GAATTCACACTGAGCAACAGGGAAGAGAAGGAACTAGAACTTGGAATGAGCCTGTGTGTGGGCCCTCATTGTTCCAGGA
TTTACAAAGGGCTACGCTGTCATCATATTAAATAAATGGCTCCTGCTCCTTGTGCCAGCTTAATTGCCACCTTGGCTCC
CGAACAAAGAAATCACAATTGAGGCAAAACATGTTTTGCTATTGAAGGCAAAACCTCCTTAGAGCAATTTTGGGGTTGT
TTTTCCATTTAATTTTTTAACTTGCAGAAATAGCCCTCTTTTTTTTCTGTTTTTTGGTTTTTTTTGTTTTCGTTTTTTTA
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Fig. 6.230

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AATGATGTACACTTGTGCTGTTAAGTTTCAGCATGGAATACAGGCTTTATGTATTTAGGCAGTAATGCTGAGTTAGATA
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CTAGGAACTTGCTGGAGATGCAAATTATCTGGTTCCATCTCAGGCCTAATGAATCAGAAACTCTAGGTGGAAGGGAGCA
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AGAAATGGTGTTTTAATTATGTAGAGAAGTAATTTCATTTTCTCAGAGAAATGGGTAGACAGATATTAGTAATTTAACT
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ACAGTAGGCCATTCACTGAGTTTGGGGTCCAGCAAGAGGAGTGAGTTTGAGGGTGAAATTATGTGTTCCATTTGGGCTA
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AGTGATTGGGCCACTACTCTCGATGTCAGTTTTAAGTGTTATGTGAATTTTAGCGTTGCAGAATTTTAGCATTGCAGAA
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TTTCATGTAGGTTGTACCAAATTTGTGGTGGTAAACATTGTATTTAGGTCATTCACAAAGAGTACAGAAATACAGATTA
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CGAGTAGCTGGGACTACAGGTGCCTGCCACCACGCCTGGCTAATTTTTTGTATTTTTAGTAGAGACAGGGTTTCATCGT
GTTAGCCAGGATGGTCTCAATCTCCTGACCTTGTGATTCACCCATCTCGGCCTCCCAAAGTGCTGGGATTACAGGCGTG
AGCCACCATGCCTGGCCCAAAAGCTAATTTTTTAAATTTTCCAGATAGAACTTCATGTGTATTTGATATATTCCATTTT
CGTGTATCCAATATTACCAAATCCATGGTATTTTTATTCTTTGTTTTAGATATTCTAATCATTTCACAGCACTTTCATT
ACACATCTGCCCAGATTTCCTGTTGCTAAATTTCAGACACTATTCTTTAAGTCTTGGAAGAGATCAACTTTTTCAGCAC
AACCCATCCCAAACAATGTAATGTTACTGTGCTATATTAAGACAACACATGTGCTGATCTTTTTCAGGCATTTCCAGAA
TTTTTATTGAATGTCAAGTATGTGCAAGGCAAAGTCCATCTAGCGATTTTAGGATTTCCAGGACCTGCCATTTATAAAC
TTCCTGGAAGAAAAATTAACCCCCTCAAAATGGGTGTTGACATCCAGTTTGCCTTTATAGCTCCTCTGATGCCTTTTAA
AAACTATATTTACCCATCTTATGTCAATAAAAAGGATGTATTTATTATAAGTTGTGGCAGATAGTGTTTTATAGACAAT
TTTATAGCCCACGTGTTAAAGGTAAAAGCAAAGAAATTATGCTACTTTGCTGTATTTTCCCAGTGATTTATGTAAGTAT
```

Fig. 6.231

TTATTCCCTGTAGGAAAAAGTTGAAACACTATTGTTGACAACAGGGCCCAAATAGGTATCAGCTTGATGTACTTTATCA
CATATTATCTTTTTGTTTTTGCATCATCCCTAGGAGGTAAATATGTAACACTTTGGTTTCAT. CATCACAACAGATT
ATTAGGCAAATCAACAGAATGAAATTAAAGTTATATTATCTAATTGAAAAAAGCCTTCACTTATATTTTCAAGCATTAG
TTCTTATCCACTTTCAATCACCTGGGGAACTTTAAAAAATCTGATGCCTGGGCCAGACACCCAGAAAAATGACATCAGA
CTTTCTGGGGACAGGACCCAGGTGACTGTATTTTTCTAAATGTCTTTCATGACTCCGCTGCTTTAGAGGTTCACAGACT
TCCACTTAAAGAACTGCTGGTTAAGAAATGCATCCCTACTTGCTCGGACTCCCAGGGATATCACTTTTCACCTGTTCCT
ATCATTTAGAAAAAAATGATTTGAATCAGCTAGATACATGATAGAGGACCCCCCCCCCACACACACACACACACCAATC
TAAGCACAAAATTATCTTGTTGTCCTCAATAGTTTGCATGGATAAGTAATGCAGAATAAGCATCATACCCAATATTAAT
AAATACTTAACAAGAAGCCCAAGATTTTACCTTGTTTCCTATTTTTTCCTTCAGCCATCACATTCAAACCTGGCCCAAA
GGTTTCTCACATTCTGGCCTCCATCACCACCACTACTGCTTTGGCCTGGCCTTTTGTCAAGCTAGAAATTGCCAGAATT
TCCTGACAACAGGCTCTGTAAGATTCAGTTTAGAGCTTACACTTTGGCTGGAGTTCATCTTCCTAAAAAGAGACTGTGA
TATGTAAATCCACAACCTGAAATCTCATTGGCTCTCCATTGCCTACCAATAACCCAAAATCCTCTACTCAGGAAGTCCT
GGATGGTGTAACTCCATTGCACACAGGTTCATCCCACAAGGCCTCTTGGTAATGGGGTACTTTCATGTGCATACTATTC
TGCTGCTCTTGTGGAAAGAGTCTTCAACCCCATTCTGTATACCC. GGACACTCACCTGAGACCCAGCTGAAACAGACCT
TGTCATCTCGGACCCTGCCCTTGGTTCTGTTGAAAGAATTAGCCACCAGATCCTTGAGGCTGTCTCAGGACTTTTGCTT
CACCTTTATTATAACAATTGGAATGCAGGCAAAATAGTGTTTCCAGTTACTAAAATAAAACTATAGACCATACAATATG
TTTGTTTTGTGCTTATTTAACTCGCAGGGGAAAATACCCATATTTTTAAGACTCTGGATTTAAAAAATCTGGTTGATTA
TATGAGAATATTGTTGGCTTGTCTGTTTGATTTTGCTTTTTTTGGGTTGCCAATAAAATTTTAGAGCAAGAAAATCAAT
CTCTGGAAAATATGTCTAAAATGCAAGAGAAATTCCAAGAAAATAAGCTAGATTTCAATTAACATGTGAAAAGCATTT
CTTAATATATACAATGTTTATTTTACATGGATAATTTATGTAAGTTATTGAATAAATAATACTTTTAGGTTTGGGTTTT
AGGGGAGGTTTTAGGTTTGGGTTGCAGCAGAGGCTTTTTCAGCTGATTCAAATATCTAGTTTTCAAAATACTCTGCCCA
CATGATTTTTAGATATGACTTAGCCTTAATTGCTGACTCAGATTTTCACCTTTTAAAAAGCAATTGTGCTTTTTCATAT
ACACTATCAAACAAGATTATAAGATATTTTAAGGTAAGAAAGTAAAAGGGCATCCACTAATGATCATCAGAATCACCTG
GAAGGGCCAATAAGCCACACATTACCAGGCCCCTTCCCCAAAATTTCTAATTCTGTAGATCTGTGGTACAGCCTCCAAT
TTTTTGCTTCTAATAAGACCCCAAGTGACACTAATGCTGCTAGTCTGGAGACCACACTTTGAGAATCACTGCATTAAAT
TAACAATTTTATTGGTTGAGTTTTCACCACAGTGTGCATATCTGTTGACTAACAATGAATAGCTTTGTTGTCGCCAAAC
ACTGTAATGTTTCCAAATATCCATATGTAGTCACAGGCCCAGAGAAACAAACCACATCTATATTAAGACTGTTTTGTTG
TTGTTTGACCCCCCCACCCTTTAGTGTTTTAATTACATCAATCCTGGAGAAAAATAACTGTTTTCTAATTTGGGCATTA
CAGATTTGCTAAGGCAATAGAAAATTCTATCTTAAACCTTTAGATTTTTAGTCTTCTAGTAAATATAAGCATATGAGTT
CCAAGCAAATTCTGAGTTTTAAAAAATTTAATAAAATTGACAAATGTTTTCAATAAAATTGGAGAGGCCCATTTCCTTG
TACAAATATTCTAAAAATAGATGATAAAAGTTCTAAGTGCCATTTTTTGAAAGATCAGCAAAATAATTCTACAAGTATT
TGTGGGCATGTTATCTTTTCTATGATACTGGTTATAAACTTAAGTTATATGAAGCAGTGTGTTAGTTAAATTATGTTC
TAGTCCTACCTTAAGGAAATAAAAACAATAAGGCTCTTAGGGCCTTTCTTCTGCACCACTGCAGTACTTGCTAACATGT
TTTAATAACATGTAGCTGAGAATGATAAACTCTAGGCAGCAGGGTGGCTTGGAGATACTAATGGAAATGCCCAGTCAAA
AAGAGCCTGGACTAGACTGAGGTCTGAAGACTCCAAATAACTTTCTGTTATCCCACCATCTTCTCAGATGGTCCAATCA
TGCTACACTCAGTCTAGGGCAATGACCCTGAGGAATGGTATGTTTGGCAAAAAAGAAACCAAAGAAGGCTACTGCCATG
CTTTAAGATTTTCCCATATATCTTTTTCAGCATCTTGAATGGGGTTTCTGAAAGTCTGAGGAGTGAACTGTGACCAAAA
TGGCTGATATTTTGAGCTCAAAGAAACTTAAATTTTTAGGTAAGGATACTAAGACCGACTTAAAAAGTATAGTATCAAT
AATTTACATTTCTTTAGCTAAGAATTTTCAGACCACTTCAGTGCAGATCACATTGACTGATTCTCTAAACCACTCTATG
AAGTGGGCAAAGTGATGGTCTCATTTTTCAAATGAGGACTAAGGTTTATTTAATTCAAGTGATTAACTCTTCAAACATC
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GGTCAGGAGTTCAAGACCAGCCTGGCCAACATGGTGAAAACCCATCTCTACTAAAAATACAGAAATTAGCTGGGCATGG
TGGTGGCAGGTGCCTGTAATCCCAGCTACTCAGGAGGCTGAGGCAGGAGAATGGCTTGAACCCAGGAGGCAGAGGTTAC
AGTGTGCTGAGATAACGCCACGGCACTCTAGCCTGGGTGAGAGTGATATTCCATCTCAAAAATCAAAACAAAAAATCT
AGTAAGGGTCTATGTCTAGCGCCTGCACTTCAAATATGTGTTTGTCATGACATCATTAGCCCAATCCAATTTTCTTTCA
TTAATTTTCTGAAATAAATATTTTTTGAGTTTATACTATATACCAGGTGCTGCTGTAAGTGCTAGGAATATAACAGTGA
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CTCCACTGCCTAATCCCTGGACTAAATGAACATTACCTCACATGCAAAAAAAGATTTTGCAGATTTGATTAGGGGCACA
GACCTTGAGATGTAGTGTCCGTATTAGCCCAGTGGGCCCAATCTAATCACACAAGTCCTTCTAAGCAAAGACTTTTCCC
TGGTTAGGTTAGAGAGAGATGTGATAACAGAAAAAGGGTCAGACAGATGCAACTTTGTTGGCTTTGGAAATGAAAGAAG
GGGGCCTGGAGCCAAGGAATGTGGACAGCTAGAAAGGGCAAGGAAGCAGATTCTCCCTTAGCCCACTCAGAAAGGAATG
CAACCCTGAAGACACCTTGATTTTAGCCCATTGAGGCCTGTGTTGGACTTCTGACCTACAGAACTATAAAATGATACAT
TTTAATTGTTAAATCACTAAGTTTCTGATAATTTGTTACACCAGCAGCAGAAAACTACAGAAAAGATAGACAATATTCA
AGAAAAATAAGTAAAAAATATATGTTGGATAGTGATAAGTGCTAAGGAAAAAGTCAGGGAATGACATAGGAAGTTATC
AGTAGGGATGCAATTTTGAATAGGGTAGCCCTCCAGGTTAACATTTGCATAGACTCAAAGGAGATGAAGGGCAAGAACA
TTTCAGATGAAGGAAACAGGAGGCTAAAAGCTGAGACTAGGACAATGATCAGCCTGACTAGATTCTGAATATCACCATT
AAAAGAGCAGTTTCTACTTCAGAATATATGTACAAGCTTCACCTGGGAACCATGTGTGATGATTTAGGATGAAGTTTCA
AAACTCCACAGATGACATTCTTATTTTTCTATAAGCAAACAGTCTTGCCCAAATAACATCATTTACTTTTTCATGTTTG.
CTGGAGAGTTCATTAAGAAATTCAATACACCGTATTCAAAGAAGATGAATTCTTACAATCACTTAGAATTTGGTCTAAG
GCAAACAATTGAGGATGTTCAGACTGAGTCTTTTTGGACCATCTACTCAACATCATTATCAACCAACAGCTTATACCCC

Fig. 6.232

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TTCAAAGGGATCTCAAATTATATTTGATCCTGAGTTGTTGCTATAATAAATTTAGAGCTGGAGAAAAATAGTCATTGTA
ATTATTTTGTTATTAGAGACCGGCTTACATATGCTAAATAAATAAAAACAAAATGAGAGTTAATTCTATGTGAAATTCA
CCATCAATTTTGATGAAATTTCCTATTGATACTTCCAAGTACAGTAGGCCTTTCGTATCCATGGGTTCCACATCTGTGT
ATTCAATCCAATTTTTTTTTTTTAATAATATGGTTGTGTATATACTGAACATGTACAGACTTTTCTTGTCATTCCCTAAA
CAGTACAGCATAATAACTATTGACATAACATTTACATTGTATTAGGTATAAGTAATCTGGAGATGCTTTAAAGTATATT
GATGTGTGTAGGTTATATACAAATACTATGTCATTTTATATAAGGAACTTGAGCATCCATGGATTTTGGTATCTGCCTG
AAGACCTGGAACCAATCCCCAAAGATCCTGAGGGACCACTGTAATCTAATTGAGGCCAAGTTTCATGGGTTTGTATCCT
CTTAAGTACCTGGGGGAAAAAACTGTCTGATTTTTTTTTTAAGTTTTAGTATACCATATATTTTAATTTTCAAAGAAAGG
AAAACAGAAATTTTATGATCTGTGTCAATAGAAGTGTTAATATATAAAGCAAAGAAATGCCTAGGTGAATATTGGCCTA
GTAATAAACATTCTGCAATGGTGCTTCATGGGAAATACGCCTTACCTTGCTACTCAAAATGTGATCCATGAACCAGCAG
TATGAATATCACCTGGACTTGGTAGAAATTCTTGGACCCCACCCTGACCTAGTGGGTAACCATCTGCATTGCAGTAAGA
TCCCAGGTGATTCATATGTGTTTTCAATGTGAGAAGCACAGCACTGGGGCCTTACCCGGGGCCCTCCTTAGGAAACACA
CCAAGTATTGCATGTTTTTTACATCCTGACAGGTCCATATGAACTCACATTTTTCCGAATACAGTGCACTAGCATATCT
ATCATATTTTGAAAAGCTATTTTCATCAAGGGAGCTTAGTATCTGGTGCTCACGAGATACCTACAGAGAAGTAGACAGT
GTTATGCTTTTGAGAATTTATCAAACCTTTTGGTCATATAATGTTTTTATTTCAAAACCTTAAATTGGTCAACATATTT
TTAAATATTTAATAGTTGAGGGATGAAGAGGATAGAAGTTTAATTAGTTCAAAGGAAGGTAAATTGGGAAGGAAACAAA
CTTTTTTGTTGAATCAGAATTTTCAGGATGAAAAAAGACACCAGTGCCCTGTGGTTCAGTGCTAACACAGCTATGATGT
GACTCTTCAAAGGTCAATGCTTGTGCCCCTATTGCTGCACATCCTTCTCCTTGTCAGTGTTGATGTGTCATTGGCTGGC
TCTGTGTCATTGGCTTGGTCTTGCTTTCACTGAACTTTGGCCTCCGTGAAACTTCTAAGTAGCATTGCTCCTACCTGTG
AATCTTGGAAACACACAGGGCAAGTGAAATTCCATTTTCTTCCTTTCTTTCTTTTTTAATGATTAATAGACAATTTAT
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CAACCGTTCAGCTTGCATCTGCCAGCCTTCCAGAGATATGTCATATTTGGCTGAGTCGAGGGTCAAGGGCAGAGTGGCC
AGGTGCGATGGAGTAAACTTTGGGAATGACATTTTGGTGACCCAGCACACAACTGCCTGAAATTCCATTTTCATGTGAC
ACTTTTGTATCTGTAACCATCAGGTCTGTTCCCTATGTTTTCTATTCCTTAACTTGGCTACCATTATACCATCACATCT
CTTTCCTTTTCTAGTGACTCCCCATTAGACTAGCTCCTATCTCATAACCTCCTTTTGTAATTTAATGGCCCAAAGATTA
GCAAATTCCAGAAATTCCTAAAGTAATCCAATTAGCTCAGAACATCATAGAGACTTTTTTTTTTCTTTGCCTCATCCTG
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TCATCCATTCTCTTAGCTGTCCTTCCCAGCTTGATGTAATAGTATTTTCCCAAACTACACCCTTTTCCATAATAAAGAT
TGTCACTCCTCCCTCAAGCTTCCTTAAGCAAGCTTTATCATATGTCTGATAGGATCTCTGTTTCTGATTGTACTCAGTA
TAGATTTGTTTCCTGGCACACATGTATGCACTATGTCACTTAATCACAGTGCCCCTATTAGATTACAAACTCCTGTCAC
CAAGGGTTTTGTTCCCTCTTTTCAAAGCAGAACTTAAAAGAATTTTGCTTACCTTTAAGCGTGCAGTTCACATCATTGA
CTTTAACTGGATGGTGGTTGATGGCTGATGGTCAACTAACAAAAAAGGAGTTGAGGAGTTGAGGTTTCAGATATAACAG
AAGGATACTTAAGACTTTTAAGGAATAAAAAACTAATTTTTCTTAGCACAATGGTTCAGATGAAAGTAAGGGTGCATCA
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TTAAAACAAATCAGAAGTAGATGACTCAGAAATGTTGACAATAAAATTATGGACAAGCCCCTTGCTACAGCAGCTCTTA
TTAATTAAATATCTGAGACAGCAGAATTAAAAGAGAAGAAAGTGATATACAATTTGGAAGGTTTGGAAAGCTGGCCATA
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AAAAGGATCTTATAAAAAGTATGTTTAGTACCATTCAATTTTTACAAATTTATGCATTTGCTTATTCATATATGTAAAT
TTAAAATTTACATATAAGCTTATACTCAGAAAAAGATAAATGTTGTAAACACACTTGCATGCACACACACACCATAACA
ACCACCATGCCATTCAGTGGTTATTTCAGGGTAAGTAGGATTGCAGAAACTTTTACATATTCACGTTTTGCCTATATTT
TCTGTTATAAGCTTTTATTTTGGCATGAGAAGAAAAAATTTTAAAATTTCCCAAAAACATTCTTTGTAATAGTCAAAA
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CCTGGTGGCTCATGCCTGTAATCCCAGCACTTTGGGAGGCCAAGGCTGGTGGATTGCTTGAGCCCAGGAGTTTGAGACC
AGCCTGGGCAATATGGCAAAACACCATATGTACAAAAAATACAAAAAAAATAGCCAAGCATGGTGGCATGCACATGTAG
TCCTAGCTACTAGGGTGGCTGAGGTGGGAGGATTGCTTGAGCCTGAGAGGCAGAGGTTGCTGTGAGCCGAGATCACGCC
ACTGCATTCTAGCCAGGGAGATAGAACAAGACCCTGTCTCAAAAAAATAAAATTAAGCAATGGGAAAAGATTCCTCTTG
CTATTTTAAATGTTGATCAACATTCAAATTTTCAGTGGTTTTAGGTAACAAGACATGTCAACCTCCTGAAACAGTAATT
GTTGTAGATAATAAGAATATATATGAGTTTGGCTTTACCTGATTTTAAAAGTGGAGAGAAAAGCTGAATTGAAGCAAGC
ATACAATTATTTTCTAAACATATCCTCTCCAGTTTCAGTGTTTTTTAATCAAACCACCAGGAATGGTGCAGCAGACAAT
GCCAGTCCATGTCTCCCCCTTCATGGTCCCTTAGTTGTCATTAATGGGTGACTGCAGAGAACCATAGGCATTTGAGGAC
TTAACAGAGATATGTTTTATATTGAGAGTGGACATAAGCTCCTCATTGCCTCATTGTGGTTTGCACTTCAAACTGCAGC
GTGTACTTTTTCTAAGCTCTGTGGTCTGCCTTCCTAGGACTGCTTGCCCTGTACTGTTATTTCAGAATGTTTATCTAGT

Fig. 6.233

GAGGAGTGCTGCATTGCTCCATTGTGCATCACAGGGCACAGGAAGCAATGGTATGACATTTTTGCTGCACATCTATTTT
TGATTATTCTCAAAAGATGAAAATGTTTATCACTGCAGGGGTGTATTAGTCCATTTTCATGCTGCTATAAAGAACTACT
TGAGATATGGTAATTTATAAAGGAAACAGGTTTCATTGACTCACAGTTCTGCATGACTGGAAAGGCTTCAGGAAACTTA
AAATCATGGTGGAATGGGAAACAAACACGTTCTTATTCACATGGCGGCAGGGGAGGGAAGTGCAGAGGAAAGGGGGAAA
AAGCCCCTTATAGAACTATCAGATCTCCTGAGAAGTCACTCACTATCATGAGAACAGCAGGGGGATCTGCTCCCATGAT
CTAATCACCTCCCATGAGGTTCCCTCCCCCAAAACATGGGGATTACAATTTGGATTACAATTCGAGATTTGGGTGAGGA
CACAGAGCCAGACCATATCATTCCACCTTTTGCCCTCCAAAATCTCATCTTTCTCACATTTCAAAACACAATTATGCCT
TCCCAAGAGTCTCCCAAAGTCTTGACTCATTTCAGCGTTAACTCAATAGTCCAAGTCCAAAGTCTCGACTGAGACAAGG
CAAGTCCCTTCTGTCTATGAGCCTGTAAAAGCAAAAGCAAGTTAATTACTTCCTAGATACAGTGGAGGCACAGGTATTG
GGTAAATACACTCATTCCAAATGGGAGAAATTGTCCAGAACAAAGGGGCAACAGGCCCCATGCAAGTCCAAAATCCAAT
AGGGCAGTCATTAAACATTAAAGTTCCAAAATGATCTCCTTTGACTCCATGTCTTACATCCAGTTCATGCTGTTGCAAG
AGTTGGGTCCCCACCACTTTGGCAGCTCCACCCCTGTGGTTTTTCAGGGTACAGTCCCCTCCTGGCTGCTTTCATGGGC
TGGCATTGAGTGTCTGCAGCTTTTCCAGGTGCGCAGTGCAAGCTGTTGGTGGAGCTACCATTCTGGGGTCTGGAGGATG
GTTGCCCTCTTCTCACAGCTCCACTATGCAGTGCCCCAGTGGGAATTCTGTGTGGGAGCTTGCACCCCACATTTTCCTA
CTGCACTGCCCCAGCAGAGGTTCTCCACGAGGGCCCCGTCCCTGCAGCAGGCTCTTGCCTGGACATCCAGGCATTTCCA
TACATCCTCTGAAATCTAGGTGGAGGTTCCCAAACCTCAGTTCTTGACTTCTGTGCACCTGCAGGCCCAACACCACGTG
TAAGCTGCTAAGCCTTGAGGCTTGGACCTTCTGAAGCAATGGCCTGAGCTGTATGTTGGCCCCTTTTAGCCATGGTTGG
GACTGAAGCAGCTGGATGCAGGGTTGCACAGAGCAGGGGGACCCTAGGCCCACCCCACAAAATAAAGTCTTTCCTCCT
AGGCCTCCAGGCCTGTAATAGGAGGGGCTGCTGTGAAGTCGTCTAACAGGCTCTAGAGACATTTTCCTCATTGTCTTGG
TGATTAACATTTGGCATCTCATTACTTATGCAAATTTCTGCAGCTGGCTGGAATTTTCCCCCAGAAAATGGGTTTTTCT
TTGCTATTACATCATCAGGCTGCAAATTTCTCAAACTTTTATGGTGTGCTTCCTTTTGAATTTTCTTCCACCAGATACT
CTAAATCATCTCTCTAATTCACAGTTCCACAGATCTCTACGGCAGGGACAAAATGCCACCAGCCTCTTTGCTAAAGCAT
AGCAAGAGTGACCTTTACTCCAGTTTCTAACAAGTTCCTCATCTCCATCTCAGACCAACTCAGCCTGGACGTCATTGTT
CGTATCACTATAAGCATTTTGGTCAAAGCCATTCAACAAGTTTCTCAGACGTCCCAAACTTTCCCACATCTTTCTGTCT
TCTGAGCCCTCCAAGTCTCTAGGAAGTTGCACATTTTCCCACATTTTCCTGTCTTCTTCTGAGGCCTCCAAACTGTTCC
AACCTCTGCCTGTTACCCAGTTCCAAAGTCGCTTCCACATTTTCAGGTATCTGCAGTAGTGCCACACTACTCTCAGTAC
CAACTGACTGTACTAATCTGTTCTCACACTGCTATAAAGAACTGCCTGAGACTGGGTAATTTGTAAAGGAAACGGGTTT
AATTGACTTACAGTTCCACATGACTGGGGAGGCTTCAGGAAACTTACAAACATGGTGGAAGGGGAAGCAAACATGTTCT
TCTTCACATGGCAGCAGGAGAGAGAAGTGCAGAGTGGAGGGGAAAAAGCCCCTTATAAAACCATTAGAGCTCCAGAGAA
GTCACTCACTATTATTAGAACAGCATGGGGGAATCTGTTCCATGATCTAATCACCTCCCATGAGGACTTTCCCCCAAAA
CGTGGGGATTACAATTTGCATTACAATTCAAGTTGAGATTTGGTTGAGGACACAGAACCAGACCATATCAAAGAGTTTG
CTTTGACAAAAGGAAGTGTATCTTTTTATTTACTTATTTACAAAGCAGCTTATTAAAGTTATTAAATAGTTTTCAAAGG
GGCAGCTTTTTTTTCTACTTTCTGTATTAGTAAGCAACCATCTATGATTGTAATACAACTGAGGTCTCCCAAGAGAGAA
TGTAGCACAAAACAAGGTGTGCTATCTAAACCTGTAAGAAATGTTGTGCAGTGAACCAACTCCAGCACAGATATGGAGC
TCTCTCAGAAACAACATAGGAAATTTAGATATGTGAAATTCAAATAGAAATAGAAAACTCAATTTAGAGTTTAGTTTGC
GTAATATCTTAGAAATGTTTTCATGGTTCAAAGCTGATATTTGACAATTGTGTTAGATCTATAAAAAATTCACAAAACAT
CCCTATAATTTTCAGATACAAAATGCTAATAAGGATTTTAAAGTTCAATGTGGACCACAGGGCTTCTGCTTTTGCAGGT
GTACCTTCATTTACATCCTTTCAGTTGAGGAAGGGTGGGTGTTTGAGAGCAAAATGTATGTATAAGGTAAGAAGAAAAG
AGAAATGAGAGAGAGAAGCAGAAGATAGTGAGCCATCAATAATGATATTAGTTGGCATTCATTTCAAAAGCCAACCACC
CCAATTTAGAGGATTTGCATTTAGGACTAATTAAATTATAAGCTAATTGAGCAGGGACTGAGTTAAGCCTACTGATAGT
GCTTCATAAATATTCATATTAACAATAATACAAGATGTTCCTTTTCAAGGCTAAAAACTTTTTCTAAATGGTGTATACA
ACTTGTGAGTCTTGGTAAGTCAATGTTGTTGCATTCCTGAATTTTTCTATCCCTTTTAAGGATAATTCTAACTCAAGTG
AACCGAAATTTTCCCTGTAGCAGTAGAGGTCCTCTGAAAATTGAGGAAGCTCTCCATGTGTAATGCTCTGAAAATGGCA
GACATTTCAGAGTCACATTCTGTATATCATTCATGTGAAATGGCATAGGCAATTTTACTCCTCAAGATTCTTTGCCCAG
AATTCGCAATTTAATAAGAACAAGTATTATGAATTGTTGAAGATTCTTCCAGCTCTCTTGGAAATAAAGGGTCTTCTCA
AATTGTAGCTTTGGTACATTAATAGTTACTCTGGGGGCCTAAATGAGTTAGTTAATGTGCAGCTTAAACATCTGAGGCA
CACCCAGAAGTAACTCAGGACTGAGGAATTCACCTTCCCTTGCTACTCAATTGCCGTTTGTGTAAAATAGTGGACAGTG
ACACTGTTTGTGTGCAGCTAGCAACTGTCTCTAAGTCTTGGGTTTGTTGGAGCATAAAGTGCACTCCAGTGCCCTGAGT
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TTTTCCTCATATTTCTATGTGGTTATTTATAGTTCAAGAACAAGTATTTAAAATATTTAAATGATAGCCATTCAATTAA
CTATTCCAAATTTTAGTTTTTTTCACCAAGGAGCTGAGCTGATTATCAAATATGCTTTATGTGAACTCTGTTTTGTTTA
AAGTGTACTGTATTTTTTAAATTTGTGAACATGGGAAATTATACAATGTTCTATAATAATTTCAAACCTGAGTTTTTTT
TAAATCTCAATGAAAGCTGTACCTTATCTGAAATGTAAATTAGTGTAAAAACCCTTTCATTCTCAATAATTGTCGGCTA
CTATCTTTTTATCTTCATGTTCATCAATAGTAAACATTCAACCTTCAACGGTTAACAAATATTAAGTGTTTACCATGTA
CCAGACACTTTTCTACATTCTGGGGTGACAGATCTAGAATTACAAGTGTATCAAATATTATAACAGACAGGTGGAAGGT
GCTACAGGATCAAATAAGAAGAGGACATAAACTTAACTGTAGAGATGGGTTGGGGAGGGAGGAAGAAGTTAGTGTTAGA
ACAGGCAACTTTTAAAGAATTAACCAGGAAGTTATGCAATTTGGGGTAAAGTGGGGAAAGAGTGTCAGGGAAATGAGGT
TGAAACAGATAATATAAGATTTGCTATTTATTCAAAAATAGTTTAGAGGCCAGTAATTGGCTGGGAGAAGAATGGTGCA
CAGATGGGCAAGTGTTGACCTGCAAGGCCAATTAGAAAGCTGTGTCTATAAGTAAAGGCAAGAGATGAGGATGACCTGG
ATTAGGAAACAAAAAAATCCTCTCTTTTACCTGCAAAAATAGCTGTTGACTTTGTCTCCCTTCCATACAAGACTTGGGG

Fig. 6.234

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TCTCCACGTTTCCAGCAAGAGACTGGGCAGTTATACATCTGTTTTTATTTTTACTCTGACCTGACTGCTTTATGCAAGA
TGGGCCAAGTCTCCCTTATACATCAGGAATGCACTGATGTAAAGACAAGATATCTGCTCCTCCAGCCACCTTTCTTCCA
GGTCTATCTGCCTTAATTCCTCTTCCCCCATTCTCAGTGGTGCATCATTCTCTACTGCTGTCCTAGTCCTGCCTCAGTT
TTCCATTAGTTTTATTTTGCAGCAGAGATTTTTTTTTTTTTAGTGATCTCCTATTGTTCCTTAATCACCCTTGAAATA
TACCTAATTAACCCAAGTAAACCTTCATGATGCATGTCTCTTCATCCTAGGAGTATAAATGATCCTTGTATATCTATAG
GAACGTTTCTATTCCAGTCTTTGAAATTGAGGAATTTCATATACATTATATATATATACACACATACAGACACACACAC
ACAACGTGTATATATGTGTATATATGTGTATATATGTATATGTGTGTGTGTATGTGTGCATATACATATTCCATTAGTT
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AGGATTTCAGAATGAAAGTTATGTAGTAAATCCTAGCCCTTTCCGAATCCTGATTTCCTAAATCTAACCTATATGGAAT
TTCTTTTTTTTTAATAAAGGTTCCATGAAACTTGGCCTCTTTCCTATCATCTTAAATCGTTCTCCTCTTATACTTCTC
TTAGTTCTACTTCTTTCACGTGAAATCCATTCTAATGTGAACCTCGGTTTCTCTGAGAGGTGCCTCACCACAATGCCCA
TGCCTTTCTGCACTTCCCACGTAGATCTTCTTGCCCCAAATACCTACTGTGAGCCTTCTTTCCTCTGTGCCTTTCATTA
TTCCACTCATCTGAAATTCAGTCGCACCAACTTTTTAAGCACCTTTATCAGGCACTGTGGTCAGTAAGGTTTACAGATG
AATAAGGCATGATTGCCAGTCACTGACAACAAATTTGGGGCAGGGAGAACAGCCCCATCCCCATAAATGGTCTCATGTT
TCTTTTGTTTGTTTGTTTGTTTTGATAGAATCTCACTCTGTTGCCCAGGCTGGAGTGCAGTGGTGTGATCTTGGCTCAC
CATAACCTCTGCCTCCCAGGTTCAAGCAATTCTCCTGCCTTAGCCTTCGAAGTAGCTGGGACTACAGGTGCTGGCCACC
ACACCCAGCTAATTTTTTTGTATTATGAGTAGAGATGAGGTTTACCATGTTGGCCAGGCTGGTCTCAAACTCCTGACCT
CAAGTGATCCCCCCACCTCAGCCTCCAAAAAAGCTGGGATTACAGGTGTGAGGTACCGTGTCCTGCCCATAAACAGCCT
CATTTATAAACCATAAATACATAAGTGCTGAGATAGAGATTTGTGTGTGTGTGTGTGTATTTGGGGGGTGGTGGATACA
CTTTGCCCAACAAATCCAATATTGTTTACTACTTCCTGTGCTTCCCACATGACTGTTCTCTCATCTTTCTCTTAAATTA
ATGGAAATACCCTCCTAACCAGTTTTCCGACTGGTGCTTTTGCCCATCCAGTCTAATCTCATAGCAGAGTTATTCTACA
AAAACAGTAATTAGTTCACATCGTGTTGCTCTCTGCTCAAAGCCATCCAATGGCATGCCATCTAGAGTCAAAGTCAAAT
TCTTGCTATGTCTGTTAAACTTCTACATGATCTATGTCTACCGCCCCTTTCAAGTGACCTGGATCTGACATCTTCTCAA
ACCACTGTCCGCCTCACCCACTCACTTCTCCACTCTGACTTCACTGGTGTTCCCTGAATGTGCCAAGCATGTTCTGCAT
CACAGCCTTTGCACTACCTGTTCCCAGTGCTGGGAATTCTCTTCCTCTAAGTATCTGGATGGCTGGTTCCATCCTGCCC
TTCAGATCATCGCTCATATGTTACTTTAGCAGTAATCCTTTCCTATTCATGCTTTTTCTTCCACCTTGACGTTCCCTGC
TGTATTTTTATTCATAGCACCTATGACCTACTGTATCCTTTGTTATCTGTTTATTGACTGCTTTCATCCAACAAGAATG
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ATAGGTGCTCAATGAATATTTGCTAAAGGATGGAAGGAAGAAGGGAAGAGGAGGAAGGAGGGAGAGAGAGAAAGAGGAA
AGTTGAAAGAAGAAAGGAAGGGAAGAAAGATGGTAGGGAAGAAGGAAGGACAGCAGAAAGTCTCATGTTGGCTTTACTA
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CAACATTATCTGGATAGAAACTAATAAGGTTGTTCTGTTTATGCTGCATTTATGGCAAGAGTTACTGACTTTCCATTTA
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Fig. 6.235

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AAAATATTCTGCAATTTCTCTTGATCACAACAAGTGTTGGAAATATCTTCTCCCGAAGAAAAGAATGATTCTTAAAAAA
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Fig. 6.23G

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TGAATCCATAGAACAGAAATTCAGTGTTTTCAGATTGCTAATCATGTATGAAAGTTTCTGAAATCATATTGTCCAACGG
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AGCAATTCTCCTGCCTCAGCCTCCTAAGTAGTTGGGTCTACAGGCACCTGCCACTACACACGGCTAATTTTTGTATTTT
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Fig. 6.237

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TTTGTCCTTTCCTTTTCACTGGAATTTCTCCATGCTTTAATTTTTATATCAAGAAAAACAATTCTTGCAATATTCCCTA
AGGAGAGTTTTACAAAAAAAGAGATAAAACAAACTACACACACACACACACACACACACACACACACACACACTA
CCTTAAGGAATAAAGAAATTATCTGAGTGGAAAATAATGGTGCTTTCAATTTTCCTAATTGAATTAAAATAATTATCTA
AGTCTTTTTGATCTTGTCCCAAAGTGAATAAAATGTCTTTTTTTAAATTTATGTTTCATTAGATATATCTCCATCTTTT
CAGTATTCACCCCAAGTTTTAATTGGGTAGAAGAATATGGAAGAAATTCCTATTGCTCAGACACCATGTTTAAAGCTTT

Fig. 6.238

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CTGAATGTGATCTAATACCTCCATTGGTTGGAGGTAATTAAACAGAGATCTGTTCTTTTAAACAACTATTTGAGTGCTT
TAAATCTGAAAGTCATGTCATAAATGTCAGACCTCTTTCTCTTTGCTTTAATCATCTCAGAAAGACTTATCTGTTAATG
GAACATCTGCATATAGGTATTTTGTTGACCACAGGTGTTCCACATAAGTTTAATCTACCTATTAGTTATATGTTCATGC
TATTGAGGACCTTTCAATAATAAACTTTACTGAGTGCTTACTCTCTTCTAAGAGTAATGCACTATGCTAGGAGACAGGA
GGGATATAAAGGTAAAGAAACCATGGCATGAATCTTTAGGGAGATTATAATTTAGTGGAGAACACTAGTGAAAGGAGGC
AAAGGAATTAATAGAACAAGACAGTTTCTCTGAGTCTCCAAAGGCACTGTAAAAATTCCTGGGGAAACTGCAAGCAAGA
GCAAAATGTTAGCTATCATAAACATTCTCATGCATCTTCAGGTTTTATCTTCATCCATAAATTTACCCATTAAATTTAG
ACATTCTACTATCTATTCCACTTGACATTCCAATTAGCAGCTCAGCCTCAGCCTATCAATATGAATGCACATCGCTTCC
CTGTTCTCTTGCTGTGCCAGTCTTTCCCCAGTGTAGTTAGGTAGTGTCTCCAATCAGTCACCATCACCATACTGTGCTG
ATTTTATATTCTGAATATTTTTTAAGCTTTTTCCTTTTTATCAATTGACTACACTAGTTTAGATCTTCATCATATTATT
GTACACAACTGCTGGGTTTTTTCTGTCTGGAAAGTCTATTTTCCTCTTTCCTCCCCAGCTGCCTCTTTCTTCTGCCTCT
GGATGCATTTCCTCATTTTCCTGCCTTCCAAGATGTATTTAATACATTCCTCTGTGCTCTCATATGCTGATACATGTCT
GTCGCTGTATTTGCTGTGTAATATCCAAATTATTTGTGCATGTCTTTGTTTTTCTCCAAAAGTTGCAAGCTCCTTAAGG
TCAGGAGCTGTTCTTTAATATATCTACTCTACCTAGTGTAGTCATGCTTAATATTTGTTGAAGTAAACAAATCTGAACC
CCAAATGTATTATGTAATCTCCATATCCCAGCACAGATAAAGGATATGCTCTTGAATTGTTTTATTGGGAGAAAACAGC
TGACAGATTAGGCTACACAACAACTAAAAACTAAGAAATGACTACATGATAAATTGTAATGAAAATCGGAAGGCAGACT
GACTGAGGAATATATCTCTAACAAGTATTCAAAGATATAATTTTGTTATTGTTAAAAAGAATACAATGAATGGAGAGAG
GGTTTGGATTTCACATTCAGGACTTCTGAAGACCAAGTATAAGAGATTCAAAAGTGATAATGTGAATCTGATTAGAAAA
TTCAATTTTGGAATTTAAATAAGAAAAAAATTAGTTGGTCTAGCAGGTGAGGCTATAATATGGAAAAATGAGAAAGGGA
TACAGCACTGAAAGTGAAAATGTGGAAAGAGAGAAGATCAGAAGGGGAGAAAAAAAGAAGAAATAAAGGGTTAGAGTTT
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TTTATTTTTATTTTAGATTCAGGGAGTACACGTGCAGGCTTGTTACATGGGTAAATTGCGTGAAGCTGAGGTTTGGGCT
TGTAATGATCCCACTGCCCACGTAGTGACATAGGACCATAGGGAGTTTTTCAACACTTGCTCACCTCCCTCCTCCCACC
CTTTTGGAATCCTCAGTTGTTCAGTGTTTCCATATTTGTGTCCGTGTTGCCTCCAGTGTTTAGCTCCCACTTATAAGTA
AGAACATGTGGTATTTGGTTTTCTGTTTCTGTTTCAATTCACTGAGAATAATGGCCTCCATTCTTTATCCATGTTGCTGC
AAGGAACATGATTTCATCTTTTTATGGCTGTATAGTATTTCCTGGTATATATGTAGCACATTTTCTTTATCCAGCCCAC
CATTGGTGGGCATATGGGTTGATTCCATGTCATTGCTCTTGTGAATAGTGCTGTGATTCACATATAAGTGCAGGTGTCT
TTTTGGTAGAATGGTTTATTTTCCTTTGGGTATACACTCAGCAGTGAGATTGCAGGGTCAAATGGTGGTTCTATTATTA
GTTCTTGAGAAATCTCCAAACTGTTTTCCACAGTGGCTGAACTAATTTGCATTCCTGCCAACAGTGTGCAAGCGTTCCC
TTTTCTCCACAGTCTCACCAACATCTGTTACTTTTTGATTTTTTTTAATATACTAGCCATTCTGAGTGGTGTGAGATAGT
ATCTCATTGGTTGAGTGTGGTTTGCTAATATTTTGTGGAGGACTTTTGCATCTGTTCATCAGAGAGATATTGGCATGTA
GGGTTTTTGTTGTTGTGTCTTTGCCAGATTTTGGTATCAGAAAGATACTGGTTTTGAATTAGGAGTTCCTCCTCCTTGA
TTTTTTTTTTTTTTTGAATACTTTCAGTAGTATCTGTATCAGCTCTTATTTGTATGTCTGGTAGAATTCAGCTGTGAAT
CCATTTGGTTCAGGGCTATTTTTGGACATTAGGTTTTTTATTACTGATTCAATTTCATTACTTGTTATTGGTCTGTTCA
GGATTTCAGTTTCTTCCTGGTTTAATCTTGGGAGGTTGTGTGTGTCCAGAAATTTATGCATCTCCTCTAGGTTTTCTAG
TTTTTGTGCATAGAGATGTACATAGTAGTCTCTGAGGATCTTTTATGTTTCTATGGGATTGGTTGTGATGTCACTTTTG
TCATTTCTGATTGTGCATATTCAGATCTTCTCTCTTTTCTTGTTAATCTAGCTAGCAGTATATCAATCTTTTTTTATCC
TTTCCAAGAACAAACTTCTCATTTTGTATATCCTTTGTATAGTTGTTATGGGTCTCAATTTCATTTAATTCTGTTCTGA
TTTTAGTTATTTCTTTTTTTTTCTGCTAGCTCTAGGTTTAGTTTTGTTCTTGTTTTTCTGTTTTTTTTTTTTTTTAGGT
GCGATTTTAGGTTGTTAATTTCAGATGTATCTTCTTGATGTAGACATTCAGTGCTAGAAACTTTCCTCTTAACGCTGCC
ATTGCTGTACTCTAGAGGTTTTGGTATGTTGTATCTCTATTTTTGTTTGTTTCAAATAACAGTTTGACTTCTGCCTTAA
TTTTTTGTTTATTTTAGAGTCATTCAGAAACAAGTTGATTAGTTTCCTGTATTTGTGTGGTTTTAAGAATTCCTCTTGC
TGTTGATTTCTGTTTTTTTTTCCACTGTGGTCTGAGAAGATGCTTGGTATGATTCTGATTTTTTGAAATGTATTAAGACT
TGCTTTATGACTATGTGTTCAATCTTGGAATATATATCATGTGCAGATGAGAAGAATGCATATTCTGTAGTTGTTGGGT
GGAGTATTCTATAGATGTCTATTAGGTCCAATTGGCCAACTATCAAATTTATTAAATACAAAATTTTTTGTTAGTTTTT
TGCCTGTCTATTGCTTTCACTCGTGTGCTGAAGTTCCCCACTATTATTGGGTGGCCATGAAAGTCTTTTCATAGGTCTA
TAAGCAATTATTTTATAAATATGTATCCTCCAATGTTGAACATATATATATTTAGGATAGTTAAGTCTTCTTATTGAAT
TGAACACTTTATCATTATGTAATGCCTTTCTTGGTCCTTTTTTACTGTTGTTGTTTAATGTCTATTTTTTCTGATATTA
GAATAGTGATCCTTGTTCTTTTTTTGTTTCCTATTTGCATGATAGATCTTTCTCTATCCATTTACTTGCAGCCTAGCCT
ATGGGTGCCATTACAAGTGAAATGGGTATCTTGAAGTTAGACTTGTTTTTAGTCTAATTTGCCACTTTTTGCCTTTTAA
ATGGAGTGTTAAAACACTATGACAGGAATAAAACCTTACATATCAATATTAACTTTGCATGGCCAAAGTTAATATTTTA
TGTTAAAACACTATGGCAAGAATAAAACCTCACATATCAATATTAATTTTGACTATGCAAAAGTTAATATTGATATGTG
AGGTTTTATTCTTGCCATAGTGTTATTATCTGGTTGCTTTGTAGTATTGATTCAGTCACTGCTTAGGGCCTGTGGGCTA
TCTGCTTGCATGTGCTTTCATGGTAACAAGGATCATCCTTTCCTTTTGTTTTCATGTTTAGAACTCTCTTAAGTATCTC
TTGTAGGTCTAGTGTGGTGGTGATGGATTCTTTTAGCAATTGCTTATCTGAGAAAGAATTTATTTCTCTTTCATTTATG
AAGTATAGTTGGTGGGATATGAAATTATTGGTTGGCATTTTTTTTCTTTAAAAATGCTAAAAATAGTCCCCCAATCTCT
TCTGGCTTGTATGGTTTCTGCTGAGAAGCCTGCTGTTAGTCTGATGGGTTTCCCTTTATAGATGATATGACTGTTTTCT
CTAGCTGCCGTTAAGTTTTTTTCTTTCACGTTGACCTTGGATAGTCTGATGATTGTGTGCCTTGAAGATGGTCATATTA
TATAGTATTCTTCCAGGAATTCTCTGGATTTCTTGTATGTGCATGTTGACTTCTCTGGCAAGATTGAGGAAATTTCCCT
GAATTATATCCTCAAATATGTGTTCCAAGTTGCTTAGTTTCTCTTATCTCAGAAATGCCAATGTCACTTTACATAACCC
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Fig. 6.239

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CATATTTATTGAAGGTTTTATTAATTGTTTAAAATTATTTTTTTCTTTATTTTTGCCTGCCTGGGTTGATTCAGAAAGC
TAGTCTTCGAGCCCTGAAATTCATTCTTCTGCTTGGTATAGTCTGTTGTTAAGGCTTCCAACTGTATTTTGAAATTCCC
ATAGTTAATTTTTCAATTCCAGAAGTTGTGTTTGGTTCTTTCTTAATATAGCTATGTTGTCTTTCAAATCTTGGATCAT
TTTTCTGGCTTCTTTGTGTTGGATTTCAACTTTCTCTTGGATCTCATTGAGTTTCTTTGCCATCTAGATTCTGAATTCT
ATATCTGTCATTTCAGACATTTCATTCTGGTTAGGGTTCATTGCTTGGGAGCTAGTGAGATTCTCTGGAGGTGGTAAAA
CACTCTGACATTTTGTATTGCCAGAGTTCTTGTGCTGGTTCCTTCTCATCTGAGAGAGCTGATGCTTTTTTCTTTTCTT
TTTTGAATTTGCTATTGTTTGGATGGAGCTTGTTGATTTTAAATTCTTTTTTTCCCTTGTGGGTATGACTGTGGTGTAAA
ATGTGTATGGGTGGATCAGCTTCATTTCTGAGTGCTTTCAGGGCGCCAAGGCTCTGTATGGGTTCCTTGGTTGCAGATA
AGTTTGTGCGGTGGCTGAGACGTTGCTTCTTGTAGTGATGTAATTTTGTTTTGTAGTGTAATTCAGGCTGCAGCCCAGT
AGGTGGCACTTAAGAGTGAGCGCCAGAAGGTAGGGGCAAGGGCAGAAGCAATGGAAAAGTCTGCAAAGTGCCCTCCTTC
AGCGCGTTTGCCTTCAGTGGGAGTGGAATTGCTGGAGAAGCCCCAAAAGTGGTCTCTTTCAGCCCACACTCTCTCGGTT
CCGCTGGGGAGAGCCACCTTCGAGTCCGCAACAGTACACTGAAGAGGGATGAGAGAGCTGAGCGATGACCCCTTCTCGA
CAACCATTCCCTGGCTTTGGTGGCGCCCCCTTCAGTGGCTAGCGCAGTGCTCCTGTTTCCTTTGACCCAACCTTTGACC
CAAGAGGGGCTTTAGCCGGCTACATCCCCCTCCCTTAGGGGCCGACTGAACCGAGGGCTAGATTTCTAGGGGAGTGGGT
CCCTTCCCTTACCACTTTTCAGAGCTGGTGGGTTACTGTCCCCCAACTGAACAAGGAAGCCGACTGGGGAACCCAGTAG
TTGATATGTAAGGGATGCTGGGAAGTGCTGGGTAAAGAAGGGTGAGTCCCTGGCGAGGGCTCCACCCCCAGGCCTGTGC
CCACGGACCTAGGTGAGGACAGGCACTCCAGCCTTTGGGCCCAAATGTTGCATTTCCCAAGACCAACCTAGCCTACCAT
GCCCCCTACCATCCTGTGCCTATAAAAACCCCAAGACCCTAGCGGGTAGAGACACAAGCAGCTGAAAGTGGAGAGGACA
TCGAGGGGAGCACGCTGGTGGAAGAGCACACCAACAGATGCTGCCACCTGGCAGGCCGTCCAGCAGAGGAATGACGCGG
AGTTTGGTCGTGGCAGTCATAGGAGAGCCCGGGCTGCTGAGCGGCTGGACTCCAGGGGGAAACCATCTCCCTTCTGGCT
CCCCCATCTGCTGATAGCTACTTCCACTCAGTAAAACCTTGTACTCATTCTCCAAGCCCAGGTGTGATTCAATTCTTCC
CTCACGCCAAGGCAAGAACCCGGGATACAGAAAGCCCTCGTCCTTGTGATAAGGTGGAGGGTCTAATTGAGCTGGTTAA
CACAAGCTGCCTATAGACGGCAAAACTGAAAGAGCCCATGGTAGCACATGCCCACTGGGGCTTTGGGAGCTGTAAACAT
CCACCCCTAGATGCTGCCGTGGGATCGACCCCCACAACCTGCATGCTCCCCTAGAGGTACGAGCAGCAGGGCACTGAAG
AAGCGAGCCACTTCTCCAGTTGCACACCTTGCAAGGGGGACAAGGGAACCTTTCTCATTTCGTAGCAACACACGCAGAC
TGGTTCCAGGTCTCAAAGCTGCCCCTAGCTGCATGTCTTGCCACCCAGAAGAAACCTGGCTTCGGGAACTCTCCTGCCC
CTGCAGTCTTCTAATGCTTTATGTCTCTTCTCTATCGCATCCAAGCATTCTCTCCTAGACTATCTGCTCAGAAGGTGTC
CACTTACTATTCTGGCGTTTCTCTGTTGGGGAGCCACACACTACCTGCTTCTGCTCAGCCATCTTGATCCCTTTCATGT
CTCTATTTATGAAGGATGAAAATGGAATGGAGACATGGAGATTGAGGATTGATGATGAGGCTTGTTGAAATTTGATGAT
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TGGACTTGCAAAGAAAAAACCTTGTTGAAGAGCAGACGTGAGCTAATGGACTAAGAGAAGCATGCAAGTGGGGAGACTA
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TATATCCAGAAGTGATCAAGGATATTTGTGAGTTTGCTTACAATATCATAAAGGCAGGACAAGACTCAAGGAAGATTCC
TAGGGAGGAAACTGGTGGGGAAGTCAAGGCCAAACTGGGCTGGATAAGGTAGAAAGAGCATATTGATGAAAATAACAGC
CTTAGGTGGGAATTGTAGGGGGTAACTTGAAGGGGATTTGAATACAGGAGCCTGTGTTAAAAACTAGCTTTTTCTTTTC
TTGCTTTTTTTCTTAGAAATGTATAGAACACATCCATTTATTCCAACACTCCAAGTGAGGAATAAGGTTTGGGCTCAGC
TTGTAGTCCGCACCAATCACTGAGTATTATTTTGAGGGGCATGTTTACGGGGAAGGTACGTTGCTGGTGGGATTGTTTC
AAAGTCTTAATGCAAGTGCCTCAGGTCTATTGCTAGACATTTAGAATTTTCTTGGTTTTTTGAAAGAGAAAGATTAGGG
ATAGTTAATAGTAGTTTAAAAGACTGTGCAACCTTAGTAACATTTCTTTGCAATAATTAATTAAAATCACAAGGAAAAT
TTTTGCAAAAGTTTGGGCATATAAAATAAATGAAAGTTATGACTAACAGTTCAACTTTCTCCTGATGTCTCACTTTTGT
GAGAACTGATGTAGACCAGAAAATAAATTTCTAGGTGTTTTTTGTTTGTTTGTTTGTTTTGTTTTGTTTTGTTTGCTTA
GGTTGACAGATCTAATTTTCATGTCAAATTAGGACACTGACTTTCCCTAACAGATGTTCAGGAGCTGAATTTGTATCCT
AATTCTAGTCTTGCTAGTATTTATTTTTGAGACCTGCAATCCATCATTATACTAACATGGCAATGCCTAGGAAGTTTAA
TGGCTATACGACTTCTAGAAAGTGCCTTTATTACATATCTGACACATTGCAAATTCACAAGTGGAGTACAAACAATTTC
AGCAGCTTAGATGACATTTTCCTTCCTGCCAGTGGTCTGAACTCACCAGGTAGCAACACCCAAAGCATTACATTCTGTC
TCCCTGAGTTGCATTGTATTTTGATATCACAGCATAAACTGTCAAAGGAAGTATTTCTCTAATTTTAATTGGGGTTAC
TAGGGTCAAAGCCAAGTTTCCAAAGCTCATTTTGTTTTTATCTTATTATCATGCATTTATAGTAAGCCATTTATTTTTT
GAAGCAATCCTTATTATAACTTGGAAAAAAACTATTTCGTAATGTGGTCAGATGTGATTAGATCACTGTTTTATGCTCA
GACGTATGTTCCACCTTTTCCCATAATGCTGTGACATTTGGGGTTTGTATAGCTTCATGAATCGAACTTAAACAGTAAT
GCATTTTCACAAATAAATAGAATCACTCTTCCTACCTATTATGATGGATCTTAAAATATCCTGCCACATCTCTCTCGTT
ATTCATTCAGTTTCTGATCTTAGTAATTTACCTATACATTAAAACATTGTAATTATGTATTTAAATCACCAGGACACCT
ATCATTTGGTCTCTAAAACTGAGTTGCTGGTTGTTAGAGAAAATGTATTTAACTTGTATTTATGAATATAACTGGATG
AGATTCTGGTTATTTAATAGGCAGTTTAACTTGAGAAACAGAGTAGATGGGAAAAATGAGAACTTACTGTGCTTTAACA
ACTTTACATTAATTTGTTGGTTCATTGTGAGCTAGTTTATTACAGTATATTGAACTGTTTACCAAATAGTAAACTCGAT
TCCAGTGTTTAGAACATGCTTTTTTAAAAATGACATTCTTAACCATTTTGCCTTATTGCAAGGCATTTACGTATATACAT
TGAGAAGCTGTATCTGATCATGTTTCTGCAGAGATGTGTTATGAACTGCTTTAGTACACGGATTTTTGAATCTTAACAA
AGGATTTTGGCCACCTGCAAAATAAGAGGTTTTCATTCTTTTTAGAATAGTCTTTTTTCTGGGTCCCCAGTGTATATTT
TGCCAAAATGCCTAAAATATTGCAATAAAAAAACTCAGAGTTTGGTACCAATCTGACAATTAATTTTTTTAAGCCTTTA
AAAGTATATAAACAATTTTAGCCAATGGACACCTCTCACTGAATTGACCATATAGATCATATTAGTCCATTTTCACACT
GCTGATAAAGACACCCTTGACACTAGGAAGAAAAAGAGGTTTAAGGGACTCAGTTCCACATGGCTGGGGAGACCTCACA
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Fig. 6.240

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ATCATGGCAAAAGGCAAGGAGGAGCAAGTCGCGTCTTATATGGATGGTGGCAGGCAAAAAGAGAGAGCTTGTGCAGGGG
AACTCCTCTTTATAAAACCATCAGATCTTGTGAGACTTATTCACTATCATGAGAACAGCACGAGAAGACCCACCCCAT
AATTCGATTGCCTCCCAGGACACATGGGAATTGTGGGAGCTACAATCCAAGATGAGATTTGGGTGGGGACACAGCCTAA
ACATATAATAGACCTATCAGTTCCCTACACTAATAAACTTCCCTGGTATTATTTCCACAGCACATGTACCCTGGAAAAT
TAGCTCCAGGTGATGAGAAATTCAGTTGAGGGCAGTAAAAAAGAAACCAGGTATTCTCACACTAATTTGAGGGTTTAAT
CTCACTGGGCCTCCTGTTTGTTTTTAGTGCATTGGAGGAAAATTAGATTTTCTGAGTAAATAGAAATGAGCAATGTTCC
TATGTATATACACTTAACTCTTCAAAAAGAACAACATAGGTCTGTACTGGCACAAGTAAAATGTTGTAGAAATTGGGTC
AGGTGCCAAGTGTGAGCAACCTCACTCTCCCTAATCAGACAGCATTTCTTATATTATTTCTGGATTAATATAAGTAGGT
AATCTTGAATAAATTCTCTAGTGAATGTTCTACATCATTTATTGTCAGTGAATGTACAGATGGGATAGTCCCAGTAAAT
TTTTTTAGAAAACCAAACCATGTTCAGGATCATGGATACTTGTATTCAAGGTTTTCATGGGCATTCAAGGAAGAAAGAG
GTGATGCAACATGCCGTAAGCCAGATTCTCTAGAAAACAGACTTGGAGGCAGAGATTTAGGCCCAAGAAATTTGTTTGG
GGGTGGGGTTCTCTTAAGATCAACACCTGTGTGCAGTAGATTTCTCTCTCTTTGGATAAAGGCAATTCCAGGAGAGAGA
CTTAGCTGAGAAGACTCCCAGCTGCTGGGGGAATGAATGCCTCAATCCTGAATGGGTATCTGGGTTGCATATTTGAGCA
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TTATCCATTATCCCTGCTATCTCACCAGGAAGAGTATAGCGATAAAGAGCAATTTCTATATCTTATACATTATATTTAA
CTATAACATTTATTGTTATTAGATAGTTAGATTAAGTAAGAAATAAGGTATTCATGATTCTTTTGAATCTGTGTGAATA
TACATATAGTTAAATAGTGACATTGAAAGTATGGTGAATTTTTATATCATCTTTCTGGAGAAATACTGAGGAAAATACT
TGAATTTTGCATAACCAAGGCATCAGTTACTATTCACAATGAGACTACAGAGAAAGCTTTCCAAAATGTTTAAATGAAA
TATGCTTTCAATGTTATGTGGTTTTCCTGTGATGTAGCTACTTAATTAGCATTGTATCAATCACCACTGACATCAAATA
TGATAACTTGCTTTACTGGAAAGTCACTGCTCAATCTCATGAATGAAATTGATGAGGTAGAAGGTAATGGTGCATTGCC
AAGAAAACAAGAAAGAAGGGAGCTAAAAATTGAGAGTTACCTACTTTGCAGTATCTGCTTTATATATGGTATCTGATTT
CAGTTTATCTGAACCCTCCCCTGGAAGTAGCTATTATCAACTTTATGTTATTTACTTACTCATTTTATTCTTTTTAAAA
AGAATAATGAGATGTTTTAAACATAGAGAAAAGTGCACAGGGCAATATAAAATCACATACATAATCATCTCCCAAGAGT
AACTAATGTCATTATAGGTAAAGTTTGCTTCATATCATTTTAAATATAAATATTGCAGATATGCTTGAAATATATGTTA
GGCCATATGTGTTCAAGCTGAGGAAAGGTAGGTTCAGAGAGAAATAATCTTGGCCAATCTCACTCTGCTAGTAAGTAGG
GCTAAGATTTGATCGTCTATACGTGGATCGTGGAACTCATGTTCTTTCCCTCCACTATGTATACTTCCTCTGTACTCCC
TGAAGACTTAATCTGGATAGAGATACATTGCCTAGAGTTTCTCTATAGTGGGAGAAATGAGCCTTTTACTTTCGTGGTA
CCACGTCAGTTGCCAATTGATTCTGGCACATAAAAATGTATTATATCTTTACCTGTTTAATTATTTTCTGTACCACATC
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ATTATGTGTATGAACATGAATTACCTGCATGGATTATTTTTAAGTTCAGAAACTGTTTTTAGTTGGTTTTCTTTCTTTT
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GCTATGGTCACCTGAAAAGAAAAGCTATTTTGCATGCAGTGGTAGGGTTGAGAAACAGCCAATATATTAAATATTTTAA
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TCCACAGTTGCTTTAGAAGAAAGAAAAAGAAATCAGTTTGTAAAATGAGATAATTGTAAGGGTATCTCATCTTTGTAG
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AATGTTGTTATTAAGGCAGTGCTGATATTATAATTTTACTATCTCTAGAGGAATACCAGAGATCTTATTAAACACATGC
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TGGGCACAAGGAAACAAATAAGGATTCTTCCATGCAGCTAAGGCACCCCAAATCTTATTAATTCAAATGTTGAGAGCAT
TGAAGCCATAAAGACAATAAGAATACTGAAACACCATGTATCTGCTTTCATTTAAATAGCCAAACAATTCTGTATTTTC
CAAGTGGGCTTCCTTAATATACATATTTTAAACCTTGTAAGGTAATTCTCCAGAGAAATATGGAGGTCTAAAAGACAAA
GTAATTCCATTTATCACCTTAAATCTCATTTAATTAGGGAATAATTCTGATATATCAAGAAAGAAACTAATTACTGTCT
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GACCACAGCAACAATGTTTTTTTTTGTTTTTTTTGTTTTTTTTTTTGAGATGGAGTCTCACTCTGTCGCCAGGCTAGA
GTGCAGTGGTGTGATCTTGGCTCACTGCAACCTCTGCCTCCAGGGTTCAAGCGATTCTCCTGCCTCAGCCTCCTGAGTA
CTACCACACTCAGCTAATTTTTGTATTTTTAGTAGAGATGGGGTTTCACCATATTGTCCAGGATGGTCTCTATCTCTTG
ACCGCGTGATCCGCCCTCCTCAGCCTCCCAAAGTGCTGGGATTACAGGCATGAGCCGTCACGCCTGGCCAACAATCCTT
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ATACAAGTCTCCTGAAATTGTGAAATATTCCAACTGTACTTGGAAAGCTCTCTGAAGATCCCCCCCTCAAAAAAAAAAA
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AAATGAATATCCTGGCTTTAAGTAACTGTTTGCTAAGCACAGTGTGCACAGCTATGCACACACTGATATCACCATTGCC
AATTTTATTAAAAGAACCACAATACAGAGCAAAGAGAAAAAGAGAAGTGTCCCTATTAATGGTATTTCAAATGCGTATT
TAAACAGCTTATCTACTCCAAACTAATTTCTGATAGATTAAATAGCTGTATTCCAAATCTTACAAGATGCACATCTCTC
CTCTATTTCTTTCTCGGCTTGCTTCAGCTGAATAGCTCAAATTGTTTGGCACTGGTAAAGCATGAAAATGTGAGATAAA
AAAGGAAACGAAGCTACAGCCTGAGTGGCCTACAGGCCTTAGAAATGGCAACACTTTAAAATTCATTATATTTTACATT
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Fig. 6.241

GCTTCAACAGTTACACATTTTTTTTTTTTTTTTTTTTTTTGGCTCTTGATGATGTCAGAAGATTTATCTGGGATGTCTGT
CTCCACGGGTAAATAGTGACCGGGGTGAAAACTGAGGGAACTAGGGAGCACCTAGACCCCAAGCCTTAGTAGTCAGCTC
TGGAATGCTCCCTGCAGTGTGGGGTGTGAGTGTGGTGTCTCTGTGTCTCTGTGTACACACTGTGTGTCTGATGTGCAGA
GGGGAAGGCAGGGAAGAAATACTATGGCAGACAGTCACTTGTGGTAGCTCTCTAAAGGGATCTGGCACTTCCTGCCATT
TCATGGAAAAGAGGAATGAGAAGGAAATAAGCATTTGTTACATGCGGGATCTCACTGGATGCTTTCAGTGCTAGAAGGG
AGCTATTTCTCCTCCTAACGTCTGTAACTCACCAGATTTCACTAATTCAGAGGTTGCTTTGAAACTCAGAGTCAAATTC
ATAGCCAGGGACTAAGTCTTCATGATGTGCCTTTTTGTTTTAGCCTCCTTTCCGGCATGTTTTTTGTGTTGCTCTCCTT
CCTGATTACTTTTTATTAGTGCTATCTTAGTTTATGTATTTTTATAAACTGCCTTAAATGCTTTCTGGAACCACTGTGC
ATTTAATCAAATAGACTAATAACTAACTAGGAAAATGATAGGGCTGATAGCATCGTCTGACCCATGAGGAAGCTGAATT
TAAGTAGCTTGTCCAAGGTCACTCAGCAAGTCCCTTGTGCTGTGAGTCTCGGTCTTGGCTTTCAGTCAGTATTTTCAAG
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CAGTTAATGTCCTATCAACTTGAAATTCCTCCCCTGCCTTTCACTCCCAGTAGAAGTTTCGCTCACCTTTCCCAGGATG
CTTGAAATACCTTCTCATGAAGCATTTCTTATCCTGAAAAAGCATCATTGTTACATCTCTTCTTAGACGCTCCTTTGCC
TTCTGTTTTCAATCTAGAAATTTGTTTCTCCCCCATCTCCATACTGACTAGGATTTTGTTTGCTTGTTTCAGTTTAGAT
CACAGGTGTTAAAAAGATGATCTTACATATTTAAATGATGGACATATGATTCCTATTATCTTCTACTTCTATGCATTCC
TGATAGACAACTCTTAGGCTGGAAACTAAAACAAGAAGGCCTATTAACTATTTTTGCAAAATAAAATTGTATTCTTAAA
GTTAAGTTCTTGATGCAGGGAGAAAACATTTTATGATTTTCATTTTACTTCCCACAATAATACAAGCATGTCAAGTATT
GACCTACAATAAATGCTTAATAAAAACCTAGTATTTTTCAGTCACTTTAACAGGGCATTGTATATTTAATGCATTAACG
CTCCATAGGCCATGTCTCCTATGTACTTTGTATGCTCAGAAAGGGAAGTAATGTACACTGGTGGTTAAGAACATATTAA
CCTGAGTACATCTTTAAACCTCAGCTCCACTCCTTGCATGTTGTATGACATTGGCGAGTCATTTAACAGCTATGGGTCT
ACATTTCTTCACCTGTAAAATGGGGATAATAATAATGTCTATTTAATAAATGTATTGTGATAGTTAACTGAAATAAAAC
ATGTATAAAGCATGCAATAGGGCACATGGGAAGTATTTTATAAGTGTTGTATGTTATTATTAACCTGTAAAGAATTCAT
ACAAGAAGGCACAGAAAAACATCAATTGCAACTAAGATGAAAAGTTTGTTGCTGTTAGAGCAGGAGTAGTTAAGATGGT
ATGTATTTCAAATATCCAAAAGCTTCTTCTATTTCTGGAGAGCAACCATGGGAAGTCTATTGTGGCATTAGATAGTGCC
ACAGAGAAAAGACAACAGGATGGTGAGGACAGGAGTTGGGAGGGTGGAGCCAAGGAATCAAATAATGTGGTTTCCTAGA
TGACATTCTTTTCAGAAGCCCAGAGGTGCCACATCTATTACAAAGTCTCTAACCAAACCATATCCATGAACCTATACAA
GATAAATTTTAACAGAGTGCAGATTGTCTTTCACTCTACATCAAACTATAGAGTCCAGAAAAAATATTTTATATCATGA
TAGTGATAATAATTGAAGGAATCAGAAGGAGTTGATCATTCTCTACAAGCTTGTTGTCATGAGGAGGGAGGTGAAAAGC
ATTAAAAAAAAGCTTAAACCCTTTTCCCAATATTTCAAAGTAGCTTTGATGGTATTTAGATAGCAAACTCATTTTAAAG
AGAATAGGATTTTCACAAAACGGAAAAATCAAGCAGCTGGCATAGTAAGTTATCTGAATTTGATCTGAATTGACGTCAG
GGAGCTGGGAATGCCCCTGGCAATCCTGCCATTCTCTCTCTTTGTGCAGTATTTTGACAGTAGTGAGAATGTCTGAGTGT
TAATCATTTTCTGACAGTGTAGGCTCTAATCTTATGCAAAATGAAAGCATTTAGAGATTTTACCTTCTTGGTTCCAATC
AGGGCATTTCTTCTAGAAAAACTATTCTGCAGTAAGATTGTGTATAACTTATTTTACATTTGTATATGTTTATATATAT
TTATATTTATATAATAAACCTAAAAATATGTAACTGCTAAGTTTGGTAGTAGCCTTTCTTCAAAGCATCATTCTTGAGA
AGAGAGTTTATTTTGTAAAACCTTACTGGTAGAGACATAAATCTAAAAAACAAAATTCAATCCAGACTTTTGTATCCAA
ATATGAATGTTAATTTCCTGGGTATATCTTTTTTTTAAAAACTTGCAGTGTGGGTATATAAGCAGGGCCAGATGTCTCTG
TGTTAATCATTTCTACTCTTTAATAAATAACGGTGGCGTACAGTGAAAATAAAGCCCAGATGGCTGACACATTGCACAA
ATCACTGGATGCACTGTATATTCATTTCATAGTTTCCTGAATACTAATAGATGATAGAGTCATCATCTTTTTATCTGTT
CCTCAGTCAAATAACAAGTAACTACTTACTGTGTCTTCTAAAAGCCACTGTGCTAGTGGCTAGGTAGGTTATAAAAACC
CACCAGATATTGCAGATAGTACCACAATATAATTTGCTTTAATCTCTATGACCTTCAGAATCTATTACATCCTAGTCTT
CCCTAGGAGACTGGGGTTTCTTGGAATTATTTTAAAAAAGTGATCCAAAACTGGATCTGATTTTGCCATGGGGATTGCT
TAACTAATTGGTTTATTAGCTTGAGTCTCATGCATCATGGTGGATATTACTGGTTGTCAAGTTGTGGTTGGTGAATCGT
AGACTTTTAGAACTCTCAGAGAAATTCAGAGTTGACCCACATTACATTTTACAGATGAGAAAATGTGAAACAAATGACT
AAATGGTTTCACTAGAGCCAGACAGTTACTGTCAGAATTGAAGCTTAAACTACAGGCTGCTAACTTGTAGTCTGCAGCT
TTTGCTAGTATGTATGTTACTTTGTCTCAGAAATGTAGTGAAATTAAATGTAGTGAGATTAAAATCGGTATCACAGGAA
TGAAGGTGCAGTAACTTTGGGTTTGCTGCTGCCATTGACTCAGGGTGGGATGAAAGAAGGGGAACGGACATCATTGCAC
CTGCTTCATTTGTTTGTTCTAGAAGGAAATAAGGAAATGAGGTCTGGCTCCACACAGTGTTTCAGTGGGGGCTTTAAC
TGGTCTAGATTATCTAGAATTTTTCAATATACAGAACTTAATAGGCACAGACACTCAAATGATAGGGGAGTCAAGTATT
CACAAAGATATGACATTTACAGAGCCAGTCTGCAAAGATAAAGATATGCTAAGTGGTTGTCTGTAGACAATTATAGTCC
ATTACAAGTTGAGCATAAAAACATAAGCATTATTCATCAGATACTGAATCACCAGTTAGTGCTGGTATTTAGATGAAAA
CATTTGGCTTGGAGGGGAAGCCCCTCAAAAACCATAGTTTTGACTTGTTTCTAAGAACCTGCCTGTCATGGAAAAGCCT
AGACCCTAGTGGAATTCCCCAGGGATGTTTTACATAGTTGGGATTCTCAGTAGTTGATGATATTGGGAAGCAGAACATA
ACCAGAGGAGCTGGGAGTCAGCCCTGCCCAGGTGGGGCCCAAGCAAAAACCATACTCTGCCTTTCCCCAAGTCACAATA
AATTGTGTTCTGCTAAGTTGTAGTTACACTGTCTCTTCAAACAGAGACCATAATAATCTTTGCTGCATTTATTCAGTTT
TATGTTTATGCTAGTGATGTTTGTCACTTAGTTAAAAATTAGCTTTAATATGGAAGAAGAGGGTATACCTCTTTCTGGAA
ATGTAACACTCAGCCTTAAAGGTAAAGGGAGCAGCAAAGGTGTCCCTAAAAAGCAACTTGCATCCTGAGCACTGTGTGG
ATAGAGGCCACTGTTGCCAGACATCCAGCCTTAGCAGGTTGCAGCCTTAGCAGGTTGTGCTGTCTTTGTATGTGGTCAT
TTTGGTACCATCCAGAACCCCCACTTACTCTCCTGCTCCTCCACCACCACCTCCAGTCCCTGGTTGCAATGAGCCACAC
AAAACAGAGGAAAGTAAACGGGCATTACCTTAGTATCAGCAATGCTGGGATATAGTTCTGCTGGTAGCTCCTCCTAAAA
ATACACCCCTAAAAAATAAAATTGTAAGAAAAAAAGAGTTAAAAATAAAAAGTTTCAGTCTCCTTGCCCTGGGCAGGTC

Fig. 6.242

CTTGGAACTGTAGAGGATTTTAAAGGCTATGAAAATTATTTTTTATGGAATTCCAGAGAGGGGGATGGGCTCAAGTTTA
TATGTGAGATACATATTTACTTATTTAATAGATTTGAAAATTTATTTCTATGTTTATTAGTCAGGGTTCTTCAAAGGAC
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TTCCAAGATCTGCAGCCAATAAGCTGGAGACCCAGGAGAGCTGCTGGTATAAGTTCCAGTCTGAAAGCTGGCAGGCTTG
ATACCCAAAAGGAGCAAATGTTTCAGTTCATGTCCAAAGGCAGGAAAAAACCAACCTCCCAGCTCAGGCAGGCAGGAGG
AATCCTCTCTTACTTGTGGAAGGTTCAGCCTTTATGTTCTATGCAGGCCTTCAGCTGATTGGATGAGGGCCATCCAAAT
GTAGGAGAGTAACCGCTTTCCTCAGTCTACCAGTTCAAATGTTACTCTTATCCAAAAACACCCTCACACACATACCCCA
GATGATGTTAATTAAATATCTGGGCACCCAGTGACCCAGTCAAATTGACACATAAAATCAATCATCCCACCATGTATAT
TTATTCATATGTAAAATAGTAGATTTATGTAGGAAATAGAAAAATGATGGTTATTATTAGAGAGGAGCATAAGAAGGAA
ATTTGTGATTTGAAGATTTAGAGTCCATTATTGCTAATGACCATTAAGCAAGAATCAAAAAATTTTGCCCTGTTATGAT
AATGATGTGTACACCATTCTAAAAATAATCCTTATTTACTAAGGCATCACAATAACTTCTATGGGTCCATTTCAGACAC
AACAAGGGCGAATCACCTTTTTCCTTTTATTTCTTTGACTTCTGTTGAATATTTTTTATGAAAAATCTTGAATTTCTAA
GGGCATATTCTTTTATCACTACAGGGATGATTTTGCCTGAATATCTGAAATAGCATTGCAAAATGATTTTCTTTAGCAC
TTTACAACTGTAAAAGAAGGTCATTTTAAAGGCAGTATTTAAAAGAACAATGCCACAATAAGTGGCTCTTCTCATTGCC
CTTACATTCTTTTAAAAAGAAGAGACCAGTAGACAATGGTTAAACCAGCATGTTTCAACAAACACATCATCACTTAATT
AATTGAAGCAACTTTCCAAAAGTATTAAGGAATTAAAAATATGTAATGCTCTAGAATAGGAGAAGCCTAATAAGCCAAAT
AAGTTCAACTATGAAAAAATATCTGTCATTTTGTTAATTCACAAGCTGGAGAGGGTATGAATGTTAAGGGGATGTTGTA
GCTAGCCTAGAGACTTACTCTTAGGTCTAAGGCAGGTACTTAAAATGTGAGAGTTGAAAAGAAATTTTTTTTTATTATT
TTTCTTTTTTCCTGAAGAAGTGGGTTTTGTAAGCACAAAGGATGTAGAGAAAAACCATGCTTTTTGATTCAGACTGCAG
TGGGGAAGGTATGCAAATTAACTTATAATTTCTATGCCAAGATTTTTTTTTGGCTCTTTCTCTAAATATTTTACTAAT
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CTGAAAAATTTTATAGGGGAAATGAGAAGAGCCAGATGATGAATGGCTTCAACAAAGGGACTGTGGTCAATAATGACAG
TGATGGGAACAACGAAATGACCAAAGGTCTTGATTTCCAAGAAAAGTCAAATCATGAAGGCTTCATGTTGGAGGAGACC
ACAGAAGTGTGGAGACTCAGGCAGGTTCCGGGATTTTAAAACTCTACCGCTGGAGCAGCCTTTCCTTACCACCACCCAA
ATGCACCCATAGAGCAGGGAATAGGGGTCAGAAATTATGGAGATTTCCACTTGAAGGGGTTCCATGGGTCTAATTCAACT
CCCATTTTAGAGATGAAGCAGGTGAGTCTCAGAGAAGTGAATGATATATTCAGGACATATGGCTAAGGGGTGACCAACT
GCAGACTAGGCTTCTTGACTCGAACTTTTTTTTTTTTTGGACTACATTGTATTACTCCCCAGTCTATTAAAAAGTTTA
ACTGCCATATTGAACAAGCAAGACTCATACAGCAAAGTCAGAAAACATTTGCCTTTCCCTCCACTGTCATTAAAAATCCA
ATAAAGAGTTAGGTCTTAGCATGCCAATGGGACTCTAAAAAGAAATCCTCTCTAGGATTTTTTGTCTCCCTTTGACCTC
TTTTAAAGGCTGTATGTGGGGTATAAGCTAACGGTCAGAATAGCTGCGCCTTTGCAAGTCCTACCAGATGTTGCAGCAA
CTCCAGAGTGTAGGAATATTTGCCATCTATTGTTTTGTCCTTGGCCTTGTTAGGACTTCTACTGCAAATAGAGACAAAG
AGCCTTCAGGAACATTCTGCTGTGCAAGTGGTCACCAAGTTCATGTGCATGACACTGCCCTGGTCATTGCTGTCCTGGT
TACTCTGCAAACACTAAAGCCGATTGGCCTTGACTTAAAATCCATTTCAAGGGAGGAGAATAACATAAATTGTTTTAAT
CAGGGCAAGTTTTATTGTAATAAAGTAAATGACACAAAGGCAGGAACTACACTGCCCTATTGGTTATTATCACCTGTCT
GGCACACAGTAACTGCTCAATAAATGAATGAGTGAACAGAAACCTATTAATTGGATCAAGTAAAAAGAGACTTATTGGA
AAGATACAAAGTAATTTAAGAAAAATGTCATATTGGAAAATAATGCCTGGCTTCATGGGTATTCCCATCCTGGCATAT
TGGACCCATTAACCTCTCAGTGAATCTCAGTGACCCAACATGGCCACATGCCATGATGATGTGACTACAGTGCCCATCG
TGAATGAAGGGAAGCAGAAGCACTCTTGAGTGGTCTGGGTTGGACCAGAACCTGGGTGTTGGGTGGTAGTGGGAAATAG
ATAAGGGAGGTAATCTGTGACATTGGCAGATGGCATTTTTATTAAATCTCAGACTATTCAGAGAGTACGTGGGAACATT
ATCTTCATTGTAATATGATCTTCTTGAAAGGATAGTTATTCAAATAAAAATATATACTTTCCATGTATATGAACAATAA
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TTTAATGAAGTTATTGAAATGCTCATGCCTATAAAATTTTTCAATATTGAAAAAAAAATTAAAAATACAACCTCCAAAAA
ATGTCTCCTGTAAGTTGGGAGACCAAGAGACCAATCTGTATGCCGTAAGATATTAGTAAAATGTTTGGAGAAAGAGCCA
AAAAAAAATGCTTGGCATAGAAATTATAAGTTAATGTGCATACTCTCCCCCACTGTAGTCTGAATTTACAAGATATATGC
CATGATTATTCTCTCAGAAGATTTTTTATGAGTTCAGTACTGTCTTGATTAAGCTTATGACTAAATAAACTTCTTTTCA
TGACATTGTCGAAGAGTATCAGTTTTCAGACAGTGAGAAAAGGACAAGCAGCTGGAATCAGCTGAGAACCCAACAATCT
ATTGTAAAAATACGGTTTATTTTACCACTCGGCATATGTATTCATACACATATATGTATATTCAACTGAGACAAAAGTT
TCACAAAATCCTGCTTACTCTTACTACATGGTGATGATCTCTGATACATGTGATGATCTCTATTCTGTTTGTTTTAAAGT
TTGTCACAACACATGAAATTTGTTTCACAACCTTCCTGTTGGTCATGAGCTCTGTTTGAAATATGCAGACTCAAGGCAA
AAATGTAAATGGGAAATTTGGAATTATTATCCAGGTTGGAAGGATTAAAAAATAGAAAATACGTATGATTTCACCTTTC
TTTCTTCTGAGGCCAGATAAAGGTAACTAGTCTGAAATCTAGTTATTTATTTATCTGGAATTGTTTAGGGTTAACCATA
GCTAGTAAGGGTAGACTGTGGATGTAGTACAGTAGCTACTTAAAAGCTGTTATGTCATGAGCCTCCCAACCATGACGGT
GCTTAGGAACTCTCCAGGGAGCATTGTAAAGTATGCAAAAGTCCTGGAAACTCTAGGCCAAGTGATTTAGAATCTCTGA
GCATGGGGCCTGAGCATCAGTATTTTTAAGAAACTCCCCAGGTAATTGTAATATGAAGCTAGGACCCAAACCATCAATA
GCAACTCTGAGTAGATGCCTTTGCCTTGTGCACTGAATACTTTTGTCATGTTAACAATGCTTTTTTGTCATATTTTATG
ACGTGTTTCTTGTGTATCACCCAAAAGCAACAGATATTTTTGAATAACAGAGGTTACTATGAGCATACAGTTATGCACA
TCATTGAACCTCAAATTTTAAGATACTTATTACTAAAATGTTATACTGTGATTTATTGAAAATTTTATGAAGAATTCAT
TGACAGGAGCAGGTTATATGTTAAGTGCTACTTTTCTAGTTGAATGTGGCTCAGGAGAAATCTAGTTAACTAAGTCAAA
TAGATAGTTTTTAATACTTATATTTATTTAAATAGTAGACTTCCATAATCCTCAGTTATTTTATGTCTTCCAAAACCAA

Fig. 6. 243

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AAATAATTTCTTTAATTTTGTCATAGATTTTAATTGAAATTGGTGAACATATGCACACATACACAAAAACTGTAGATTT
AGTGTCTGTGGTATGTAATTGATGAACTGTTTTCTAAAAGGATTTTGAACGTTTAGAAAACTAATTAAGATATTTAAAT
TATTGACAAATAGAGGATCTTTCTGTACCCAATTTAAAAAGAAGATACAATGATTTGTGGTTAGTGTATCCATTTTTCA
AACATAAACAGAAACTTCAGAAGTTCTTAAAAACAGGTTGTCCTTTTAGATAATTTCTACATTTCCTCTGAAATCTTAT
TTCTGAATTAAGTCTAGATGTTTACAATCTATCTGAAAAAAGTAATGTTGTTATAAACCTCAAATCCGTGCTCATACAA
ATTCTTACCATTTCCAATTAGATAATACTCTGGCAAGAATTACTAATGCCTGAAAAAATAGATGTAAATATTCCCATAT
GATGGTAATTATTAAATGTTATCAACTAAGTATAGTGTTTTGTTTTAATTTTAACTTTTATTTTAGATACAGGGGGCAC
ATGTGCAGGTTTGTTACATGGGTATATTGTATGATGCTGAGATTTGGAGTATGGACTCTGTCACCCAGATAGTGAGCAT
AGCATCCAATAAGTAGTTTTCCCACCCAAACTCTCCTCCCTCCCTTCCCTTCTAGTAGTCTGCAGTATCTGTTGTTCC
CATGTTTCCGTCCATGTGTACTCGGTGTTTAGCTCCCACTTACAAGTGAGAGCGTGTGGTATTTGGTTTTATGTTCCAG
CATTAATTTGCTTAGGATTATGGCCTCCAGTTGCATCCATGTTGCTGCAGAGGACATGATTTTGCTCTTTTTTATGGCT
GCATAGTATTTCATGGTGTATTTGTACCACATTTTCTTTATCCAAACCACCACTGATAGGGTGGGTTTCATGTCTTTGC
TATTGTGAACAGTGTGGAGATGAACATATGAGTGCATGTGTCTTTTTGGTGGAATGATTTGTATTCTTTGGGCATATAC
CCAGTAATGGGATTGCTAGGTTGAATGGTACCTCTGTTTTAAGTTTTTGAGAAATCTCCAAACTACTTTTCACAGTGG
CTGAAGTAATTTACATGTCCACCAACAGCGTATAAGCAGTCAATTTCCTCCACAATCCTGCCAGTATCTGTTGTTTTTT
AACTTTTTAATAATAGCCTAAATGTAGTATTTTAAAATATCTAAGAGTTTCTTACCTGAACCTAGCCAATGTATTTTTT
CATCTTATTAAATTACATACACTTTTATTTTAGCCCTCATTGATTTATTAGGTATATAAGTAAGCATATTTTGTAGAGA
TTCTTTAATTTTTTTTTTTTTTTTTTTATTATACTCTAAGTTTTAGGGTACATGTGCACATTGTGCAGGTTAGTTAAT
ATTTTATAGCACTAATATGTCCTTTTGTTTTTTAAGAATAAAATTTATAAATTGTTTTATAAGAGTAAAATCAATGTAA
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CTCATTTTAATTTTTTGTAGGTGGTTTTTACAGTTGCCTTTATATATTCTGTCAATTGTGAATAGAAATTCAAATTCTT
GAATTTGATAATTGTGTCATTTTTGTGTTCCCTAGCTTTTATTTGTAGACTGTTGGTCCTATTGTTCTTTTTCATGTGT
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CGTGCTGTAATGCTATTAATAAGTGTAGTTTGCAAGCAATGTTTCAAAATTTTGTCTACTTAAAATTTAGAACAGGATA
TTTATTTTGTAATTTTTGGATAGATAGAAAATCTCCTTTCTGCCATCATGAAACTGATAAATGAGTTATAGGGATGGGA
GGAGATGGACATTTGACCACATGCAATGCAGTTTTCCAAAGTATATAACCATAATCTATATTAAAGGTAAATGCAGTGT
CAGATGGGCATTACTTTATGCAAGATTGTGTTGCTCATGAACTTACCAGCCAAAGGAGACTATCTCTCACTAAGGCTTA
TGTAAAACTGCCAGGAGAGAATAATTACAGTGGGAGTAAATTTGAACACATGCAGTAGAATGTCTAAAAGTTAAGCTTC
TCAGAGAAAATGTACATATGCTTGGATTTTACAAATGATGTATTTCCAGGTATTCACTAGGCCTGCTAAAATAAGTTAG
CATTTTTTAAAGATAAATATGTTACTGAAATATATCCTACATAAAGTACCAAATTATAGGTTTACAGTTTGATGAATTC
CTGCAAAGTGAACACATCTATGTAATCTCCATCCTGATCAGGAAATAGAACATTGTCAATTCTCCCAGAAGTCCCTGGT
ACCATACCCCAGTCCCCTCCCACCACATCACATAAGATTAGGTTTTGTCTGTTTTGTTTTTTTACCATTTTGTGTTCTT
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CTTCATAAAATAGAGGGAAGGGAGAATAGGTTGGTGGGAGGAAACTCGTGCTTCTGATTCTTTCCCTAGCCTATGATCA
ACATAATCATGCCTTAAGAACCTTGCTTTCTGTCTTTGAGTGACATGCTTGGAAAAGTTTTTGAAAACATCAGTTTGTG
TTCTTTACATATTCATTGTATAACTCTATTTTATATGGTATTAATAATAAAATGGAAACAAACAGAATTGTGGAGGGGA
AAAAAAGAGAAATTTTAAAGACAAGCCTGCCCTCTAATTTCCTTTTTGCCATGAGTAGTCAGTGGGAAGACTCTGGGAG
CTTGGGGACATTGAGGCATACAGCAGTCAACAAGCCTGAGCCCTACAACGACTGAGTCTGGAAAGTTTAGATTCTGAAA
GCCTAAAAAAGCACTGTGTAAGAGGCTAAGCCAACCTTTTTGTTGTTGTTAATCTTTTTACTGGGAGACAAGAATGGAC
CTTAGGGATAGGGTAAAAAACTGAGAGCCCCCAGCCCCCTTTAAGATATTTTCTCTAAACCTCTTAAAGTTAACTCTGAT
TAATTTCTTGGTTAGAATTCAAGTTATGGTTCCTGCTCTGCAAAACTAATAGGTAGTTTAAGGTGGGAAAAGTAAAAAT
TCTTGCCATCTCCTACATTGACTTGAGAGTGAAAGTAAGACTACAGATTTGTTATCCCCAAGACAGTTGTAGAAAACTG
AGATTATGCTTTATAATACCCTATTTCAAGTGGGGTTAATATATAACCATGGTTCTTAGCCTTATTTGGGGTTAGAGAC
CCCTTTGAGAATCTGAATAAAGCTATATGTCTTCTCTGAAAGGTGTACCTAGGTACACATAATTTAGCCACTGGCCTTG
TCCGTTTGCAGGCCCCGTAAGCTAATGCACAGTTCCCTCTGAGGATCCCTCAGCTTTATTCTAAAGTCTAAAATTCTCT
AACGGTTGCCAGTGGAAATAATCCAGGAGTGATAGAAGTATGGTCCTTATTACAGTTATCTAGGAAGGAGACCACACTG
TCTTCATCCTTTACCCTTCCTTGGTTTCACCTTCTTCCTTCATGACTTGAAGATTCCTTTATTTTCTCTTACTTCCAAC
TCAAAATTATATTGGTTCATGCTGGTTTCCTTTGTTTAACAGAAATATTCCTTTTGCAAAATGGTAAACACCTTTTTAG
GGGGAAAAAAAACCCAACCAACACTGTTCCTACAGTTGTAGATAGTAATTAAGTCAAACCCTTAGTTGGTTTTCATTTT
TGAATTTAATACTGCAAAAAATTAACTTTTTTGCATAAAACTTTTTATGACTTCAGTTATTAATATTTTATTATTTTGA
TTAATTTTATGTAAGTCAAGATAGTCTAGGTTATTCTGCAGTAATCCAAAATCTAAGTGGCTTAATATAACAAATTTTA
TTTCTTGTTCTCTTCATACATACCCAAGAGGGATTGGCAAGAGGCTCTACTTATTGTAGTAAACCTGGGACCCAAGATG
ATTGATGCTTTATCTCAACTCAGGCTTCTTCACCAGGGAAAAATGATGTAATGAATCCGTCACAGGTTGCCTGAACTTT
CACCTAGAAGTGACACATACCACTTTTGCTTAAAATAATTGGTCAAAGTATATCCCATGTTCATGTAGGATACCAAAGG
GGAGGCTAAGGGCAATCTTACCATGTGACCAGAAAAAGGAAAGCTAGAAGTATTTGGTAGACACATTCACTCTTTTCTG
TACTCCTTTCCAAATTCTCCTAACTTTTTGGAGATTTGATTGATGTTAAGCAGAAAAAGAGGGTTTGAGTCTTATTTTT
GGTGAAAGAAAATATTCTTGGTTAACTTTGTCATATAATATATCTGAACTTAAAATAAGTACACAGTGGCATATAATGA
ACTTAAAAGAAATAAAATACTAGCAGGAATGTAAAAAACCTGAACATAACACTGAATTGCTCTTGTTAGTGTTATGCCA
CTTAGTTCCATCTACAGTCTAGCCCTTGTAATAACACACACACCGTGATTTTAGTTAAATAGGAACATGCCCCTTTCTG
TTTCATTTAAAATTCTAGTCAACACAAATTCTTCTAGAAAAATGAATTAAGAATGAGTACTATTCCCAGAGTACATTCT
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Fig. 6.244

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GGTTTGTCTATAAATTCTAAGCTAAGCTTTATATGGGAACTGTCAAAGGCACAGTTGCAATGTTGCTGCCTTTTTTATT
CAGGAGAACTGACTTCTGTTGTCCCACTTGGACTGATATGTTGAGTGATGAAAAGCTGGGCAAATCTCCTCAGTCACTT
TCTTCTGATAGTTTACTTTATGGCTTCCTTAGTGGTCTGCATCATTCTTTCAGCCTGACCATTTGCTTGAGGGTGATGA
AGAGTAATGATGACAGGTTAGATGAGCTCTAGCTTCACAAATTTCTTAAATGCATCTGAAACAAATGCAGATTCATTGT
CAGGAATAATAACATCTGGGAATCTGGACATTGAAAATCATTGGTACAGGGATGTTCTTGTTGAAGTAGAGACACTATA
AGGCATTTTGAGATGGTAGCCACAACAAACAAGAAATTCTCCTCCTTGAAAAATAAAAGAATTATTGATCAAATCTTCC
TGATCATTTTTTATAGGAAGGAAAAGTACCTTTGGTGGATTATATGGAGTGGCTTGGTGTTGTCTACAGTCTTCTAAAT
GACATAATTAAGTACAGGCTTTAGATATAATTATTAGTGTATATATCATTTTAAAATTCCCAGGTGAATTGTAAACATT
ATTGTAATAATGTGATAAGTTATGTGATTTCTCTATTCTGGCCACTCGAAATTAAGAATTAAAGCTTTGATTAGAAATT
AGAGTCTAAACTATATATTTGAGAGGCATTAGTATAGTTGAAACTATGAGAGTATTTTGGTTTACTGAGCTAGTTTATG
TGAGAAAGCAAAAGACCTGAAACAGTCTTTTTTCAGTATGTATTGTATTCTAGACTTTTTTCTAGGTCATTCTTTCTTT
CCATATTTTCTCTTTTTTTTTTTTTTTTTTTCATGAGAAAGCAGAATGATTCAAACTGGGTATAACTATATTAAAGGC
CATGAACCTAGCTTTGTCTGACAATAAAGTCATGCTCTTATTGCTAGATTAAACTTCCTCCTGTTTATAGTGTCTGAT
GGGATTAAAGAGAGAGGATCTTATGTAAGTAAGAAAGTAATCTTTTTCCTACAATCTGCTTTTAGCCCTTACTTTCCTC
CTATGTTGAGAGCATTTCTAGTCCAAAGTTCTTTTTTTTAAGTCATTAAGATGTATTATTCAATCTCAAGTAACTTTTA
ATCCCTTTGTAAAAATATTAGCTCTTTAATTAGGCTTGGTGTGCCTAGCCAGCTGTCTCCAGCATAATTTTGGGTAGGC
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GAGACTTATATGACCTGTCAAATCAATTCATAAAAATTCCTATTTTGAAGGGGGAAATTAAAAGTCAAGGTAATGGAAATG
TTATATTCGTATATTCGCAGGAGATGAATGACATGGATAGGATGTCTTTGAAAGTTAATGAGAAAGCTACTTGCCAGGC
AAGTTGTTCAAGAGCAATGTTAGCAAATGAAATGGGTTGCTGCCTTTGGGGAAATGATTTATGATGATGAAAACCATCT
GGGTTTAGGAGGTTGGTTGATGTGACATACCTCTCTTCCAGCCTTTCCACTCTCTCTTCTTTCCTGTGCCCAGCATAAA
ATCAGTCTTTGCGTATAAAGAGCGCTATGCAAGTTGGCTTCCAGTCATACGAATATGTGGTATGATTTCAAATAAGATT
GTCATTGGATGCTTTGGTCAGCAAGCATATTTACTTTAGATTTGTATTTACTCTTTGTAGTTGAACTGGCTAAACTCTA
ACAAAGTATGCTTCATTTGACTTTAATAAAATATTATATTTCCAAGACCGTATGTAGTAATAGCAGAAGGTGTGATTTC
TCTTTTTGCATAAATTGGCTCTTCCCTTCTCTTTCTCTGTGTTTTTCACATCTCCCTACTGTCATATATATTCTGACCT
GCCTCTGGTAGGGCTATTTATGTCATATCCATAATTGTTATATATTAACCATTTGAATGTGGAATAAGGCAAGTACAGT
TTAATTTTTTTATATAAAAAATATGGAATATACTGTAAATGAGTTCTCTTAATCACGGAGTAGCTACCTCTGGAAGCTG
TAACTGTTGGTCTTGGGCTGTAATATATTATACTACAACCTTAAAAATGAAGAAAACAAAAAGACAACACTCTATGTGC
TGCTAGACAAAGGCCACAGTTAAAGCCAAACTGCTTGACACTTTATATTCTTCAGGCCTTTTTAACGACCTTCCCTTTG
GGTGATGCATTTTGTAGCTGTTGTCACTTGTACAGCAGAGAGGGAAAAGGAAATGAATAATCTGATTTGAAGGGGTGTA
GAGGGATGTGAGTCTTAGGAAGATCCTTAGGATTTACAGAAGCTATAAAGCTATTTGTGCTGCTTTGTGGTGCTTTGCA
CCCTGTGTCAGTTAATAGTTTATCTTTACCCCATTTTATGCTCTAAGATATAGATGATGTTTTAAAAATCATTCTCATA
GTCTTCTTTCCTCAAGCTCCACCCCAAGTTTTCTAGCTCCTGACCAGCACTGTTAGCTTTTTTGATTTAATCTTCACAG
TCTTCTCATTTATGAAATACAAAATGAATGCTATGTCAAGACAGACTTGGAAAAATCTTTTAAAATGGCAATACAGGCT
CACCTGCAATTAAAGTCATTCCATTCCATTTAGTGTCTTTTTCTTTTTCTTTTTGAGACAGGGCCTCACTCTATTGCCC
AGGCTGGAGTGCAGTGGCAGGATCATGGCTCACTGTAGCCTTGACTTCCTGGGCTCAAGCAGTCCTCCTGCCTCTGCCT
CCCTAGTAGCTATGACCACAGATGCATGCCACCATTCCTGGCTATGTATTTTTTAATTTTTAGTAGAGACAGGGTCTCA
ATATGTGGCCTAACCTGGTCTCAAACTGCTGGGCTTAAGCAATCCTCCCACCTCAGCCTCCCAGAGTGCTGGGATTTTG
GGATCCCAGGTATGAGCCATTGCACTCGGCTCCATTTAGCTTCTGATGAGTGTAAATATTAACTTTGAGGTCAGTTCAC
AGTTTGCTGATATTAGGACTGATTTTTTTTAAGTTATATTTCATTTAATAGTGGAGCACCTTAGGAAAAAACAAATCTC
TTTATATATAGAATTAAAATTTGTGAACTTTTCTGAAGTACATTTTTATCCTTTTAAATATGCCACAACATCTATCCTGT
AAGTAAAATAAATATTTCTAAATGTCTAAAGCAGAGAAATATTATAAAATTAATTAAAATATTTCCTGAAAAACAAAGTA
AATCAATTTCTCAATTTACATTATTCAGATAAAAATAGCTCAAGAACTAGCTAAACAGTATTAGTGAAATTGAAGGTCA
AGAGATGCTAGAAATTAGTCAACTCAGGTTTATGGAAGATCTTACTATATACTCTGTAATATTCAAAACTATTTCTCGG
CTTAATTCACTGATGAAATTATGGTATTTTGGTAGTTGGAATTACCAACTGTCTGCAACAGCATGAACTGACAAGAAAA
TAGTGCCCTCTATGAGTAGGAGTACAGAATGTTATCTTATTAACAAGAACCAGGGGACAGTAGGAATGGACAACAGCAT
TGGTTAGTGGCACAGACCAATTCAAATAATATATTTTGTGTGATTTTTTTTTTCCTTACCAGCAGTTTGTTGGTGAGGC
AGAATTAAATAGGGAAGAGAAATGCACATTCCCTATCTCTGTCCCTGCAGTTTCATTAAATTCTGACCAAATTATGTGA
CTCAGCATTCTCAGAGAATAGTCTCTGACAGGTAGCACTTACAGGAGACAGACCCACACCTTCTAGCATGTGATTCTTA
AAACTCTGTGTGCAGACTGGTGCCCAGCTAACCACATAAAATTTACCAGGGGTGCTTATTAAAAATGGAAATTCCAAAA
TCTTACCTCCAGAGATGCTATCTTGGTGGGCCCTGAGAACATCATGTTTTTGACGAAGTCGCCCAATGCCTTTCTGGCG
TGGTCAAAAGCAAGAAGCACAATGCCTTTTTGTGCTTCTTGGTTTTGACCAAGTCACCCAGATGATTGTGGTGCAATGC
CTTTTTGTGCTTCTGCCTTTAGCGTTGGTTTGCCCAGCCAGCACCAGATCAGGCAGCTCTATTAGCTGCTGCTATCAGC
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TTTGTTAGCTCTTGCTTGTCAAGAATGGAATGGGGAAATCATTTCCTATGGAGGAAGTTTTTCTTTGGCTTTTGGTAAC
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TGTTCAAGGCCATCCCCTGCCTTCCTTGTGTAGAGGGCTTTCTGTGACCAGCAGACAGCAGTCAAAGACATCGTATCTATAC
ACAGATGGCACATTGATGAAAACTACTGATGTTTGCCACTTATTATTATTCTTTTTTCCTAAATATTTACAAAAAATTA
CTTTTTCTCTTTTGTATGCCACATCTGACTATGTGTGTTAGTTTTTGTTTTTTGTTTGTTTGTTTGTTTGTTTTTTGAGA
TGGAGTCTCACTCTGTCAACCAGGCTGGAGTGCGGTGGCGTGATCTCAGCTCACTGCAAGGTCCGTCTCCTTGGTTCAT
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Fig. 6.24S

GCCATTCTCCTGCCTCAACCTCCCAAGTAGCTGGGACTACAGGCACCCGCTACCACGCCCTGCTAATTTTTTTGTATTT
TTAGTAGAGACAGAGTTTCACTGTGTTAGCCAGGACGGTCTCGATCTCCTGACCTCGTGATCCACCCACCTCTGCCTCC
CAAAGTGCTGGGATTACAAGCATGAGCCACCGCACCTGGCCGTGTGTTAGTTTTTATATCTATGTTAGTACCGCAAAAA
TGTCTAAGAAAGCAGACCCTTCTACCAACACTACAATGTCTCATATTGCAGGAGGCCTCATAGTTAGGAAGACAACTGT
TAGCAAAGCCCTTGTTTCTCTCATGAGCCACCAGTGTCTGGCATATCTCAATGCACTCTTCATGATGCTCTAAGCTCTA
AGTTAGGCCTATGGTTGCATTGAAGCTCTAAGTTAGGCCTATGATTAAAGTCTTCTGGCTCATAATGATAAAAGCCATT
ATTGTAGGCAGTTAGAACCTGTTGAAGGACAGAGACGTGATGGATTACAGTCTGAGATAATGTAAGTTGTTTAAAAAAG
TGAAAATAAAAGAAAATCAAAACTTTGCTTTACCTATTCATTTTTAAAATAACCAAGGCATACCCCTTTTGCTGTCTTA
AGTTTCAGACAAGGGTGCAACTTCTACAGTCATTCCTTGGTGGCTTTACATGCTTTATTTGCCATGGGCCATTAGGTGC
TATGGGTGTTTTCATCCCCACTTTGGTGCTTTGTCAAGACATCAAGCATCTCCATGAATATACTTTAATCTTTCCCTTT
TTTGTGTTTTGTTTTTTAGGTAAGTACTTTTGGTATTCTTCTTCCTTAATGCTGAAAGAAAATGCAGCTGCCTGATGTA
GAACAAATGGCTTTTGAAAGCAAAATCCAAAAGATAAAAAATAATGTGAACAGTAAAGAATGACACCATACCAGATACT
GGTAAGAATTTTAAGTGGCATTCAAACACCCCTCTCTTTTTGGAGAGAGGACTAACAGTACAGGAAGATGCGGGGAGGG
TGGGGAGGGGATTGTTGCCGTGCAAGTAAAAAGAAAATTACAGACAGTGTTCATGAAATATTCATTTACACCTGCTTATT
GGTTAGATGAACAGCCTTTGGTTCTGAGCTGGCTCACCATTCCCAGGCTGGAAATTATTCACTTAAATGCAGCTTTTAA
ACATTTTCTTAAAATCCTCAATGGAAACAGATGTGCACATGCTGAGGGAACTTTTTTTTGATCTTTTTCTAGAAGAGGG
TTAAATTAGAAAAAAAAACCAGTATTTTTAATATTTAATTTCATGTGTATAGCTATGAAGCTATATACTTAAATGCTTT
GTAACATATGATCATGAATATATGTATAAACTTCACCCAGAAAAATCCATAAAGCTTTTGACAATAAATGTGTATAATC
TGGATAAATGAAATACCAGTGAACTCAAACATAATTTATTTCATGTTTTATGTCTTCTCTTTCTCATTTTCCATGTACA
ATTCCTGTGTACATATGAAATAACATACAAAATACAATTGTACGTATGCTTCAATGGCATTTTGCCCAAATTCATAGTC
CTTTTTTATATGGCCGTGGATTCGAGATTTAAACATATATGGTAAATAGTATGTTTCTGTTAACTTCAGCATAGCCCT
TTTAGTAACAAGACATTGTTTACTATAAATTGAGATTATCTATCCAATGTGAGTAAATATATTATTAAAAATACATTAA
CATTTAAGATGAACAAATAGCAAAGGTTAAAAGGTTTCAGAGACCATTACCTGTTACTAAATCATTTCTCTCTCCCATT
TACTAGTCTTATCTCATTGAGTTTCCTGCTGTCCTAGAATGTCATCCCTCTGTCTCTGCTTTATTTAGTTTTTGTCTAT
ATTCAACTCAGACCCTATATCATGTACAAAGCTTTCCTCAAGTCTTTGAACACAGACTGGTTTATTCCTGCCCTAAAAT
CATTTTTTAATAGAATCCATGTGTTAATTTTCTGAGGCTGCCATAACAAAGTGCCACAGACTGCAAGCGTAAATAGTAG
GAATTGATTTACTGATAGTTCTGGAGGCTGGAAGTCTGAAGTCAAGAGAGCAGGGTTGGTTTCTTCTGAGGCCTCTCTC
CTCAGCTTGTAGATGGCTGACTTCTCTCTATTCCTTCACGGGGTCTTTCCTCTGTATGTGTCTGTGTCCTTAGCTCCTC
TTATAACAACGTCACATTTGATTAGGGCCCACACTTATGACCTCAACTTAATCACCTCCTTTAAACCCTGTCTCCAAAT
ATAGTCATATTCTGAGGTACGAGGACTTCAACGTACAAATTTTGGAGGTATATGTACGAGGACTTCAATGTATGAATTT
TGGGTATAGGATTCACCCCGTAACAGCCCATGTTAGTTTCTCACTTCATTATTCTCTGATTCTTCCATGTATATTGGTT
CTATCTTCCCTAACAAGTTCATCAAGGATAAGGATTGATCATATTTCAAGTTTTGTTTACTTCATTCACAGTTCCTAGA
GGGCTGGCATATAATATTAAATGCTACATATTTGTTGTTTGAAATTACTGGAAAAAGTAGCATAACAGGACCAAATGAG
AGCCTCTTTTTCCAATTATTGCTGACTTTAGCTCCCTTATCTGGACAGAAAAAAACAGTAGCAGAGGTTTGAGTCAGGC
AGTGGTCAAATGAACTGTCTCTGCTTTCTTTCTGGGAGGCCCAACAGCATTCTCGTCAGCAGGAGCATTCTGGCGAAAG
GAAATGCTGATCTCTGCAAATGGGCAAAAGTGTAAAGAGCATTGAACCCAGCCTCATACCACAGAAAACAATTTGGGCT
TGTGGAAAATAATTACCCAGTGAAGAGTCTTCTGAACTAGAGGCAGGTTTTTGGAGGGGGAAATCATAAACACATTTGT
GGTTGCAAGGTAGCAGATTGGGTGAGCTGAAACTAAAACAAATTTTGGTTTAAGAAATTGTATTTTAAGTTCTGAAGTC
ATACTTAATTCTTAATAATTCTAATTGTACACACGCTAATTTAAATGGAAGATGTTTACTTCATTAAAATTTCAACACT
AAAATGCTGGTTCAGTTTGTTGTATTACTTTCTTCAGGTTTTGTTATATTCTATTTTTCATAATTTCTAGAGCCAGCAT
TCAAAAGTAATTTCTGTGCTTCAAAATAGGATCATCCTAAATTCAAATTATATGTAAATGTCATTGAATGCAGATATAA
ACATGGTGCCACAATTTTATAGAAAATTCTAAAAAAACAAAACAAAACAGAATAACCATGTTGCTCTCGTACCAAACGT
GTCTTTGGGTGGGTTACTAAGTCCACCTGAGTCTGAGATTCTCCATCTGTAAAAAAGGAATCAGAATACCTACTTTGGG
CAGTTGCTGATATTAAATTTAAATAAGCAAAGCAGTTAGCCCAGTCATGTCAAAATAGTTGAGCCCAATGAATGGTCTT
TTTCTTTATCTCTCTTCAGGTTGAATTTTGCACAGTAGATAATCCATCTATCACAGAGGTGTGCAGATGTGACTGATTC
ATTGATTGAAACATTCATTGCCCCATTTAAACAATGTTTAGGTATTATTATCCACTGTATCCTGGTTTCATGCTGGATG
TTGGGTAAATAATGATGAGAAGAAACAGACAACTCCAACTTTAATGGAGAAATTGACATTAAAAGTCTAAAAATATGTA
TAAAATTTTAAAAAATCAGTGTTGTGTAGTAAATACACAGTACCAGGAGATTTTATAACAGGGAGGGTATGTGTAAA
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GGAGAAGCAAGAGAATGAAATAAATAAAAAGATGAGTGTCTGGGAACAGCATGTGCAAAGTCCTGATGTAGGAAACAGT
ACTGAATGGACAGTACGGTTTACCAAGGGAGGGCTGGTGAGGCTGAGATGAAGAGAGCTAACAGGAAGCCAGTAACTTG
AGATGAGGCTAGAGAGGAGAGCAGAGCCGAGCAAAACCTTATAGACATATTAGGTCTTCTACCCTAAGAACAA
TGGGAAGCCACCTAATTTTAAGAGAAAGCTTAATGGATGTGTGTGATTCAGTGCAGGGACTGCTACCGCTCTTAATCCT
CGAAATATCCTGCTAACCAGGTTGGCTTTAGTGGGTGGAAAAGTGAGGCCCATACATATTCAAATTTGTATTTTGAAAA
GATTTTTTTTTTGTCCTCCAATGCTTTTGCCAGAATATAGAAATTCAAAGTATGTGTATTTATAGCAAATACGTAGATG
TTTTTCACCTTATAGGTGTGTTTTTATCACATTATTTTTGCACATATGATTCCTGCATGACCCGAGTGAAGGGGTGGGT
TGCCCCTCCACACCTGTGGGTGTTTCTCATTAGGTGGAACGAGAGACTTGGAAAAGAAAAAGACACAGAGACAAAGTAT
AGAGAAAGAAATAAGGGGACCCAGGGAACCAGCGTTCAGCGTATGGAGGATCCCGCCAGCCTCTGAGTTCCCTTAGTAT
TTATTGATCATTCGTGGGTGTTTCTCTGAGAGGGGGATGTGTCAGGGTCACAAGGCAATAGTGGGGAGAGGGTCAGCAG

Fig. 6. 246

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ACAAACACGTGAACAAAGGTCTTTGCATCATAGACAAGGTAAAGAATCAAGTGCTGTGCTTTTAGATATGCATACACAT
AAACATCTCAATGCTTTACAAAGCAGTATTGCTGCCCGCATGTCCCACCTCCAGCCTTAAGGCGGTTTTTCCCTATCTC
AGTAGATGGAACGTACAATCGGGTTTTATAGTGAGACATTCCATTGCCCAGGGGAGGGGGCAGGAGACAGATGCCTTCCTC
TTGTCTCAACTGCAAGAGGCATGCCTTCCTCTTATACTAATCCTCCTCAGCACAGACCCTTTACGGGTGTCGGGCTGGG
GGACGGTCAGGTCTTTCCCTTCCCACGAGGCCATATTTCAGACTATCACATGGGGAGAAACCTTGAACAATACCTGGCT
TTCCTAGGCAGAGGTCCCTGCAGCCTTCCGCAGTTTTTGTGTCCCTGGGTACTTGAGATTAGGGAGTGGTGATGACTCT
TAAGGAGCATGCTGCCTTCAAGCATCTGTTTAACAAAGCACATCTTGCACAACCCTTAATCCATTCAACCCTGAGTTTG
ACACAGCACATATTTCAGAGAGCACGGGGTTGGGGGTAAGGTCACAGATTAACAGAATCTCAAGGCAGAAGAATTTTTC
TTAGTGCAGAACAAAATGGAGTCTCCTATGTCTACTTCTTTCTACACAGACACAGTGACAATCTGATCTCTCTTGCTTT
TCCCCACACCCGAGATCTTTTAATATTCTAACATACTCTTATGTTCTAAATCAGTGACACTTAGAAAGAAATGTTGACT
TGAACATCCAAGTACAGGTGCTAATTTGACCTAGCAGAAACATTTTTTAAGGAAATCTCTCTGCACTGAGCACTTGCAT
TATCTAGCAATGGAAAATTCAAACAATAGAATGATTATCATAAATTCCTTTGAATACCTTTGTTAGCAGAGATGAAACC
TTGGCCACCTGGCTTTAAGGGAAAGCTTAATGAGCATGTGATTCAGTGCGGGGACTGTTACCTCTCTTAATCCTAGAAA
CATCATGGGAACCAGGTTTGCATCAGTAATGAATTTTTATAATGTGAATCTTTGATTTTCTTTCGCTGGTAAGAAACCT
TATTTGGAGATGCTATGTATGTGAGTTTAATCTCATTTTCCATTCCTTCTTGAAGCTTTCTAGGTTAACCACCTGAAAC
AGTGGTTACTATAATCAGTAAAGATTTATAACCAAAGTATAAGGAATTGGGGAATTTCCATATGGTGTGTCCTCACAAA
CAAACGTCATGTTTCAATATGATATGAATGCATCTCCATCCAAATTATTATTTTAACTGTATATTTTTCCCTCAGTCAG
CAATTTATGTAACACATATATTTCATTTCATCCAACTCCTAATAAGAGACCCCTAAATTAACCTTGAACTATGATTTAC
ACCTTCTCAATATACAGTCAATAATTGAATTTAATGACTAACCAAGGACATTTTAGTCATTGCAACTGCTTACAAGATT
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GAGCCAAGCACACACCTGGGGTATCTTGGTACTACCTAGAGACCCTAGCTAATTTTGCCAGGGTGGATTGGTAGAATCCA
AGGAAATAGTTTTGCAAGTGACACAATTGGTGGATGATATAATAAGATAATGAAGACTAAAATAATTTGAAGAAGGGAA
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TTATAAAATCACTGCTTTGATGCATTAATTTGATCATTCTAAACAGGTGATGAATATTGTCTTATGTTATTTGCCTCCA
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AATTTCTTAATTACCAAATGACTGTGTTTTATAGTGTCCCTTAGGAAACCAAGTTTTAAAACTGTGTCTAAAGAGAACC
TGATTTAATTTTTTGATGTTTCTTTTATTTTTGCTTGATTATATATTATAATCATTTTGTAAAAAAAAATCCAAACACT
ACAGAAATATAACTTGGATGTTTATAATCTTCGCTTCTCTCTTAGATAACTATTTCTAAGAGTTTTTGTATACATATTT
AGATTTTTGATACATGTAAATATATATAATCTTTTTTACTAAAATGGTAATTTACAACATGCTTTTAAACAAACTTTTT
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CCCCAAGAGGACTCTTAGATATTTGAAAGTAGCCCATAATGGTAAAGTAAATTCATTCCTGACTTTGCTTGAGGAAACT
AACATCCTCTCCATCCTGCCTTGAGGACAAGTGTGTTCCTTCATGGCTGACACATATCTTGGCAGAGCACATGTGCTGC
AACAGCACACAGTATGGGGTAGGAACAGGATTTGAGTGATTTACTAATCAAGTGTTGGCTAGAAGAGAGGAGACTTGTA
CCACCCCGACCCTGCATCATGCCTCTTGCTCTCCTCTTTAGGGTGAATGTAAAGGGGAATCTGCCCAGTTGGGGTACAG
ATAAACCTTTTGGTTACCAGACTGTTGGGGTTTAGATGTCAACTCTTTATAGGAATTGTCTAGGAAAGTCAGCTGTCCA
GGCTTGGAGACCTTAAGGAAGCATGGAAAGCTGCAGCTCCCTTTTTCCAGGAAAAGGCAGTCACCCGTCTTCTTGGAAG
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GTAACAAATGTGATATCTTGACATCATCTGGTGGTTGGGTGACAACTGTTTAATTTTCTAACACAGTTGGTTTCTGCAA
TGCCTTTTCCCAAGTAATTTAGTATTGTCTTATCATTAAAAAATTTGTTATACTTTTATTAAAATACTTTCACAGTAAAGTTTTA
ATCCATTTTCCTATGGATTTTTCATTTTGTAATATTTCTTTCCCATTTATAATCATATAACATATACATTTAATAAAAG
TATATACAATATATATCAAGCATATATGGTATATATACTATAATTTTGTCTGTTTTCTACTTTCTTTCTACTTATGTCT
ATTTTCTGTATTTTTTTCCATATGGACATCATTTTTCACATTTAATGTCTTCTTAACATTACAGGGCAGTTCTAATTGTT
AGAAGGCCTCCTTATACTGATGGCTAGTTTGACTCATTATAAAGTCAGTTATTGTTCTTGGTTTTATGTCTTAACAAGG
CATCCCCTCCTCGAACCTTAGAGGAATTGATATCTGACTTCACATGCTCTAGGGGATGCCATGTAAAATTTTGTAGAAA
TGTTTAGTATTCTGGCATGTGTGTGTCTCTAACTTTGTCAGATGATTAGAGATCTGTGATTAGTTAACAACCACACAGGAC
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Fig. 6.247

TGAATTAGCTTCCTTCCGTAACTAAAAGAATTGAACAGATTGAGTAAATTAAGTGGATGCCTCATTAGCTCCACAAAGT
TTATTAGAATTAGAGTCAAAATCTAATACAGGGCTTTGGACTACTAATATCTGCTAGCTACCTTTGGGATGATTTCAGT
ATCATAGGATCATCTGACCAAACAAGCTGTCTCA__ATTCCTGTTAGAGTGAAACCTTGGCTGGACATTAGCTTCACCTG
GGTAATATTAAAATACTGATACCTGGCCCCAGCCTGGAGAGATTTTGATTCTATTAAATTGCCAGAGTTAGGGTTCAAG
GCAAGGTACATCCAGAGTTGACAACCACTGGATTTGAAGCCAAAATGTATAGAATGTGAAACAGGTAAGCACTGTTGGA
ACTACCGAGATAGAGACAAACACATTCCAGGCACTAGCCTCTGCTTCTGTCTTTTCTGATGTAAGAATATTTCGCCTCT
CAGAAAGTTGCTTCCCAAATGAATACTTTTAAAAATTAACCTGGTTTTTTTCAGTACATAAGACAGGGTAAGGAGAAAAA
GTTCACCTGGTTTTTAAAAACACTATTGTTTAAACTTTAACAGAATTATCTTCTCAAAATACTTAGAAATGGAGTAAAT
GTTTCTGCTTTGATAACACTGAAACCAAAGCTAGAGAGTACAGTTAAAAGGGCCATTAAAAACAGTTTTTATTCTATAA
AAATAAATTAATCATCGAATATTCATTATAAATTCAATCATGAAGGAAAATACACATTAAATTATTTATACTAAGATAA
ATATAAACCTCTATTGGCAAACAACTTCGTAGTTAATTTTTCTAATTTACCACTTTCTGCATCTCATGCAAATTATGTC
TTCTTGCATTGCCAATAAAATATGAGATTGGGGAAGGATGCTGCAATATCCATGAGAAAGTTTCATGTAAGTGCAAGCA
ATCAGATTTCATTGCAGATCCAGAGTAGTAACACTGACAGAACTATGTCAGTTAAGCTTTCTGCATTATTTTTATGCGT
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GGGTTTGTTTTTATCCCCTTGCTTAACCAATCGTACAGTTAATTCCCAAAATGTGTTATTTTTTAACATGCAAAGCAGT
TCAACAACGATAGAAGGTGTAGGGATTCCAGCCACAGACAGAGCCTGAGAGAACAAACAAAGCAGATTCAGAAAGCAGG
CAAGTAGCGCCTACTTTGTTTCAGTGGTATGTCCTCTGCAGTTCAAAATCCTTCAGTATCTCTTAATTAATTCATAGAC
TATTTCCAGTGAGGTTTCACTGAGAATTCCTTAATGTCAAGAGCTCTCTCTGTATCTTAAGCAGTTAGTGTCATTGAAA
TTTACTTATGCAATGATTCTTTCACCAAATATGTATTGAAGGCTTATGATGTGCAAGACATGTCCCAAGTTCTGGTAAT
ACAATGATGAACAAAATTAATAGTTTCTTTCTACACAGTTTTTGGTTTATCAAATTTATTACATGTTTAATAAATGTTC
ATTAAGTGCATAAATAGTAAAAGAATCAAAGATGTGATTTCAGCACTATGTATCAAAACCACAGAATGGCTGAGTGTTC
TTTTTAAATCCATGTATCATAGTGTATTAGAAGGCAAAATAATAAGAAAGCAGATCAGAAGTCAATATGGTCCAGTAAC
AAGAATAAGGAGTCAGATTAGAGTCACATTAACGGTTCACATTTTCCTAGCTGTGGGACTGTAGGTCGTTATTTTCTTG
TCTTATTTTGTCTTTCTGTTTTTCTTTCTTTCTTTTCTTTTTTTTTAGACAGAGTTTCACTCTTGTTGCCCAG
GCTGGAATGCAATGGCACAAACTCAGCTCACTGCAAACTCCGTTTCCTGGGTTCAAGCGATTTTCCTGCCTCAGCCTCC
CAAAGTAGCTGGGATTACAAGCGCCCACCACCATGCCCAGCCATTTTTATTTTGTATTTTTACTTGAGATGGGGGTTTCA
CCATGTTGGCCAGGCTGGTCTTGAGCTTCTGACCTCAGGTGATCCACCTGCCTCGGCCTCCCAAAGTGCTGGGATTACA
GACGTGAATTCTATAGGTCTCATTTGCACCATCTGCAAATGTGAACAATGGTATTGTGTTCATTTACCCATTCAGCCAA
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GCCTTCAGCAAGTAGTAATCTTTTTGCTGGTGGGAAGTCTTGCCTTGACGTTGATGGCTACTGATTGATCAGGGTGGCG
GTTGCTGAAGGTCAGAGAAGCTGTAGCAATTTCTTAAAACCAGGCAACAATAAAGTTTACCACATTTATTGACTTTTCT
CTTCATAAAAGACTTCTCTGCAACATGTAATATACTGTTTGATAGGATTTTACCGAATGTAGAACTTTCAAAATTGGAG
ACAATCCTCTGAAACCCAACTGCTGCTGTATCAGCTAAGTTTATATAATATTTTAAATCTTTTATTGTCATTTCAACAA
TTTTTATAGCATCTTTACCATTAGTAGATTTCATCTTGAGAAACCACTTTCTTTGCTCATCTATAAGAAGTAACTGCTC
ATCTCTTTGTTGCTGAGATTGCAGCAATTCAGTCACATCTTCAGGCTCCACTTCCAATGCTTGTTCTCTTGCTATTTCT
ACCATATCTGCAGTTAATTCCTTCACTTGAACCCTTCAAAGTTATCCATGAGGGTTGGAATCAACTTTTCTCAAACTCC
TGGTAATGTTGATATTTTGACCTCCTCCCATGAATCATGAATGTTCTTAATGGCATCTAGAATGGTGACTCCTATCCAG
ATTTTTAATTTTCTTTATCCATATACATCAGAAGAATCACTGTAGAAAGTAGCCTTCCCAAATGTATTTCTTAAATAAT
AAGACTTGAAATACTCTTTGAATCATGGCTGCAGAAAGAATGTTGTGTTAAGCATAAAAACAACATGAATCTTGTGTAT
TGTATTAGTCTGTTTTCATGCTGTTGATAAAGAAATACCCGAGACTGGGAAGAAAAAGAGGTTTAATGGACTCAGCTCC
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AGAGAGAGCTTGTGCAGGGAAACTCCCATTTTTAAAGCCATCAGATCTTATGAGACTCATTTACTATCAGGAGAATAGC
ACAGGAAAGAGCTGCCCCCATAATTGAATCACCTCCCACTGGGTTTCCCCCACGACACATGGGAATTGTTGGAGTTATA
ATTCAGGATGAGATTTGGATGGGGACACAGCCAAACCATATCATTCCACCCCTGGTACCTCCCAAATCTCATATCCTCA
CATTTCAAAACCAATCATGCCTCCCCAACAGTCTCCCAAAGTCTTAACTCATTTCAGCATTATCTCAAAAGTCCACAGT
CCTACATCTCATCTGAGACAAGGGGAGTCCCTTCTACCTATGAGCCTGTAAAATCAAAAGTAAGTTAGTTACTTCCTAG
ATACAATGGGGGTACAGGCATTGGGTAAATACCATTCCAAATGGAAGAAATTGGCCAAAACAAAGGGGCCATAGGCCCC
ATGCAAGTCCAAAATCCAGCAGGGCAGTCAAATCTTAAAGCTCCAAAATGATCTCCTTTAACTCCATGTCTCACATTTG
GGTCATGTTAACGCAAAGGGTGGGTTCCCATGGTCTTGGGCGGCTCCACCCCTGTGGCTCTGCAGGGTGCAGCCTCCTT
CCTGGCTGTTTTCACAGGCTGGTGTTGAGTGTCTGCTGCTTTTCCACACATGGCGCAAACTGTCAGTGGACCTACCATT
CTGGCATCTGGAGGATGGTAGCCCTCTTCTAACAGCTCCACTAGACAGTGCCCTAGTAGGGACTCTGTGTGGGGCTCC
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GCATCCTGGCATTTCCATACATCCTATGAAATCTAGGCAGAGGTTCCCAAACCTCAATTCTTGACTTCTGTGCACCTGC
AGGCTCAACACCACATGGAAGTTGCCAAGCTTGGGGTTTGCACCCTCTGAATCCATGGCCTGAGCTGTACCTTGGCCAC
TTTTAGTCACAGCTGAAGTGGCTGGGACACAGGGCACCAAGTCCCTAGCCTGTACACAGCATGGTGACCCTGGGCCTGA
CCCATGAAACCATTTTTTTCCTCCTAGGCCTCCAGGCCTGTGATGAGAGGGGCTGCCATGAAGACCTCAGACATGCTCT
GGAGACATTTTCCTCGTTGTCTTGGGGTTAACAATTGGTTCCCCGTTACTTGTGCAGATTTCTGCAGCTGGCTTGAATT
TCTCCACAGAAAATGGGATTTTCTTTTCTATCGCATTGTCAGGCTGCAAATTTTCCAAACTTTTGTGCTCTGCTTCCCT

Fig. 6.248

TATAAAACTGAATGCCTTTAACGGCACCCAAGTCACCTCTTGAATGCTTTGCTGCTTAGAAATTTCTTCCACCAAATAC
CCTAAGTCACCTCCCTCAAGTTCAAAGTTCCACAGATCTCTAGGGCAGGGGTAAAATGCCAGCAGTCTGTTTGCTAAAA
CATAACAAGAGTCACTTTTGCTCCAGTTCCCAAAAAGTTCCTCATCTCCATCTGAGACCACCTCAGCCTGGACCTTATT
GTTCATATCACTGTCAGCATTTTTGTCAAAGCCACTCAACATGTCTCTAGGAAGTTTCAAACTTTCCCACATTTTCCTG
TCTTCTTCTAAGCCCTCCTAACTGTTCCAACCTCTGCCTGTTATCCAGTTCCAAAGTCACTTCCACACTTTCAGGTATC
TTTTCGGCACTGCACCACTCTACTGGTACCAGTTTACTCTATTAGTCTGTTTTCATGCTCCTGATAAAGACATACCTGA
GACTGGGGAGAAAAAGAGGTTTAATAGGCTTACAATTCCACAGGGCTGGGGAGGCCTCACAATCATGGCGGAAGGCTAG
GAGGAGCAAGTCACGTCTTACATGGATGGCGGCAGGCAAAGAGAGAGCTTATGCAGGGAAACTTCTGTTTTTAAAACCA
TCAGATTTTGTAAGACTCATTTACTATCATGAGAATGCTGCAGGAAAGACCTGTCCCCCATAATTCAATCACCTCCTAC
CAGGTTTCTCCCACGACATGTGGGAATTGTGGGAGTTACAATCAAGATGAGATTTGGGTGGCGACACAGCCAAACCATA
TCATGTATCCATCAGAACTCTTGGATGACCAGGTGCACTGTCAATGAGCAGTAATATATTTTAAAAATCTTTATATAA
GAGCAGTAGTTCTCAACAGTGTGCTTAAAATATCTAGTAAATCATGCTGTCAACAGATGTGCTGCCATCTAGGCTTTGC
TGTTTCAGGCACAAGCACAGGCAGAATAGATTTGGTGTAATTCTGAATGGCCCCAGGATTATTAGACTGGTAAATGAAC
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ACATTGACTTATTCTTTCTAGTTGTGAAAGTGCTAAATGGCATCTTCTTCCAATAAAAGTTTGTTTTGTCTACATTTGA
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TGCATTTTTGTGTTGTGGAGATGGCATTTTCCTTAAACCTCATGAGACAGTCTCTGCTGGCTTCAACCTTTTCTTCTGC
AGCTTCCTCATCTCTATCAATCTTTATAGAATTGAAGAGAGTTAAGGCCTTGTTCTGAATTAGGCTTTTCTTAAGGGAA
TGTTGTGGCTGGTTTGATCTTCTATCCAGACTACTCAAACTTTCTCCACTTCAGCAACAAGGCTGTTTCAATTTCTTAT
CATTTGTATTCACTGGAGTAGCACTTTTTATTTCCTTCAAGAATATTACTTTTGCATTCACAACTTGGCTGTTTGGTAC
AAGAGGCCTAGCTTTCTGTCTGTCTTGACTTTTGACATGCCTTACTAAGCAAGCTTAATCTTTTCTAGCTTTTTATTTA
AGTAAGAGATGTGCTGCTCTTCACCTCACTTGAACACTTAGAGGTCATTGTATTAATTGACCTAATCTCAATATTGCT
GTGTCTCAGGGATGGGGAGGACTGAGGAGAAAGAGAGAGATAGGGAATGGCCAGTTGGTGGAGCAGTCAGAACACCCAT
TAATTGAGTTTGCCATCTTATATAGGCATGGCTTATGGTACTCCTAAACAATTACAATAGTAACATCAACAATCAGCAA
TTACAGATCACCATAATAGGTATTAAAATAATTTTAAAAGTTTGAAATATTGCAAGAATTATCAAAATGTGATACAGAG
ACATGAAGTGAGCACATGCTGTTAGAAAAATGGTACCCATAGACTTGCTCTACACAAGGTTGCCACAAACCATTTTGTT
AAAAAAAAAAAAAAAGCCATCTCTGCAAAGCACAACAAAACAAAGCACAATAAAATGAGATATGCCTGTTTCAGTCACT
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TTAGGAGTAAAGACATTAAATGTGATGAATGTTTTTGTGTTGGCCTTTTACCTATTTATTCTCAGACCACATATTGCCC
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ATGAAGGTTTACTGAGAAAGTAAAAATTTAGGCTAAAACCTCAAGTGTGAATATAAATGTATCAATGGAGAGGGGAAGAA
TATTCTAGATCAGAAGTGTCCAATCTTCTGTCTTCCCTGGGCCACAATGGAAGAATTGTCTTGGGCTACACAGAAAATA
CACTAACAATAGCTGATGAACTAAAAAAAAGAATCACAAACAAAAAATCTCAATGTTTTAAGAAAGTTTACCAGTTTGT.
GTTGGGCCACATTCAAGGCCGTCCTGGGCCTCATGGATTGGACAAGGTTTTTCTAGATCATGTGCAATGGGAAGGAAAA
TGAGGCATATTTTAGAAACTAAAGATGTAGACAACTGGAATGTGACATACAAGAGAAAGAAGGTCAAGAATGTTTGAGG
GCTGTCAAGCTCATTGGAAGACATTACAGTGGAATGACATGGGCATTTGTGTTTTTTAAAGTCAGTGGCTGCATTGTAT
AGAATTACTTGGGATGCAAGCAGGGAACCATTTAGAAGATTATTACAGACAGCGGTGTTGGTATGGAGCAGAATGATAG
CAGAGGCAATGGAGGAAAGTGAGCTTATCTGAGAGATAGAGGTAGGAGGTGGAGTCTGCATGTCTTGGTAATTGAGAGA
CTCTGTGAAACAAGATTGAGGAAAGGGTTAGGATTATTTTCAGGATTCAAAACAGAGCAACTGGGTAGCTGAGGTTCTA
ATCACAGTGAGAAGGAATATTGGAGAAGGAAGAGTTCCATGGTTGAACTTCAGGTATATGTCGGTTATCCCCAGGGAGA
GCTGTCCAGGAGTCAGGTGCGCTGTATGTGTCTGGAGCTCAGGAGGCAGATGCAGACCAAAGAAATTGATGGGGAGCTA
TCCATCTACAGATGGTCATAGAAGTTATTTGAGTCAATAAAATCAACTAAAGGGGGTGCAAGATTAGGAGAACAAGAGA
GTCCTGTCTCAAGCCTAAGAACTGCCAATATTTGAAGACAAAGCTGAAATGAAAGACATGAACAGAGACTTAAGAAGAA
AACCAAGAAAGTTAGGTGCTATGGAAGCCAAAGGAAAGGAGTGTTTCAGAAAGGAGGGAGTGGCCAATAGCTTCAGAAA
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CAGAGTGAGGAGGAGGTAAGAAGTATCTGGATAGGGAAGTAGGAATTGAGAGGGATCTTATTTTGTACAGTGAAAAATA
AGTGCTTATGGAAAAGATTTGGTAAAGAGGAGGGAAAAGAGTAAGGGAAAAATGGATGAAGCTGCGTATCAGAGAATGTG
AGAAGGACAAAGAACCAAAGCACAGGTGGAAGGATTTGCCCGAGTTGGTAACAGGAGGGAAAATGGAAATGATGACTGT
GGGAGTAGATGTGTTTGTCATTTTGATGGCTGGACATGGAGAATGTTCCTTTTTGATGGTTACTATTTTCTCTTTGAGT
AGGAATATCATCTGCTGTGTGTGGGAAGTGGGGACTCAAGTATTTGAGAAGAAAGAAGAAGGACGTTTGCTGTAGTCTT
TGCAGAAAATGAAAGTGATTTGATTAATAGTTGTTAGAATTTAAAAATGTGTGCACAGTGTTTGGGTTAAAGTTGTTTT
TAAAAAACCAGCCACCTTGCCTTTTATTCTTTATGCTGTGAAACCTCTTTAGAGCACTCAGTCACCTTTTGGCCACTAG
ATGGACACAGTGTACTCAGTGCTAAACTGCTGACCCACCAGGTTCCTTTTGTTACCAGCCAATACTGACAGAGTGATTG
AGTTGTGAACTTATTGTTATGAATTGAAACTTATTTTAGAGAAAAAATCTTAAATAACACATTCCATCTTACTTATAGT

Fig. 6 249

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TCATAGGTCAGAGAAAGACTGTGGTGCTACAAAAACATTAGCCAATATATTATTTGCTTTCACGCTAAGTGTAATGTGT
GTAACATGCTATCTCTTTGAAATTTTTTGCCTTAAAAATGCTAATCAGTTGGCACAAGGCGATCATTTACATAGTCAGA
ATAGAGCTTTTGGTTTAGCATTTTATCTTAAAATAAGGCAGAAATGGCATTGCTCTGGATGTCAGTATGGTGCATTATA
ACCCAAGTGGTGGAAAAATAACTGCTAAATGGCAAACACATAGAAACTGAATTCTGCTAGTCAGCTTCCATTTGGTAGA
GATATGTGTGCCCTTGGGTAGCTGCAATGTTAGCTATTATTAATAGTTCAAATCTTTGCTTCATAAAAGTTCTGCATAT
AGTGTTGTACAAATTGAAGTGATTCAAGGAAATCATAGTTCTGTGGAGCTTCCTCCCTGTTTTGTAGTGGAGATTGGGA
ATGGGGGTGGACCATAAAGTAGGTGGTTTTTTTTTTTCCTGCCACTCTTAACTAATTACACACCCTGCCATATCCCCCAC
CAACATAAGACTTCAGACTGAGAAAACCTACATAATTTAACCAATGTTAGAATATAGGCATTTTTAACGTGCTGAAAAC
AGTGTTGTTCTTTTGGTTCCTAAACTAAAATCATTTGATACATTTCTCTGTCCTATGAAGGAAGCCATGAATGCTATGA
GTAAAGATATTTGTGGCAGGTAAGAAGAGGGAGTTAAGTAAGGGAGTTTAATATGAAAAATTCATATAAAGGCTCCAAG
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GAAGGAATCGATTTGGGAGAAAAACTATATCTGCTTAATGTGAGGGAGCACTGTGGAAAATTTCCAGCACAAACTTTTC
CTGGGCCTCAACATTTCTTGGTAATGAACCTAGATTTGGAGTGATCAAATAATTTGCTGTCTCAAGTAGTAAAAGGGGG
ACTATTACTAACTTCACAAGAAGACAGGCATAAACCCAGCAACATGTACAAAGCAGGATGATCTCAGGTAACTCAGGAA
GATGCTGGAACACTCTGGGTGAAGGGCATTAATGCTGTTCACAGTGGAACACAGGACTGCTAATGGTATTTTTATGCTT
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CCCTAGCTGCTGGGAATGGTGGCTCACTCCTGTAATCCCAGTGCTTTGGGAGGCCAAGGCGGGAGGATCACTTGAGCCC
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ACATTATTCTACATTCTTCCTTTTTACTCTCCATATTGATACAGAAACAAAACAAAACACATTTGAGACTTACCACTGC
AAAAGAGTGTTCTTTGAATTGTGGCTGCAATTATTTCTGTTGCAAACTTTAGTGTATTATTGTTATAATGATACTTGAA
GTAATACCCAGATGTATTGAAAATTCAAAAGACTTCATTGCCAGCTCAACCTGCACTCTCAAGCACTGCTTACTAAGGG
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TTGTTTCAGCTTATTTTGACTAATTAGGCTGAGGTTACAAACCACCCTGATATTTCAGTTGCTACAGCCACAGTCTTAT
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GTTTACTTGTCTCTCTCCTTCCCTCAGTGGTGGGCCCTTGGAGGGCAGGAGCAATGTGATATTCATCCATGAGACCCAA
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CTTGAAGAATAACTAGAAGGTTACCAGATGAGAAAGTACAGGGAGGTAAATCAGGGAAGAACAGCTGATAGATTCAGTT
GTAGCCATGTTGAGTGAAGGTAAGATTCTTGGATTTTGCTTCATGAACTCGAAAGTCTACATAACATCAATCTTGGCCA
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TCTAAACAGAGCGTGAACCTCTGACATTGACTCTGGAAATTACACACATTTTGTTTTACTATTTTAAAAACACACATGA
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CAGGTATGTATGACCATGGGGCAGACAAAAGACTGCTTTAAAAATCACTGATGTGTATCATGCTTTTTCTTCTTCTTTA
GTATTTCTAGCCATCAAAAAGAAATTATATTGTGAGGCAGATGGCAGTAATATATTTTTCAAAAGTGAATTATATCCTG
TTTCCCCATATAGGTAGGTAAATAAATAAATAAATAAATTCTATATAAGCAACATGACATAATTCCCAGACTATGAATT
TCTGCTAATCTGAAGCATTTGATTAAGACTGGATTTAAACTATGCTTACAAGTTGCCGGAAACTACTCAAAAAGGCTAC
ATTCTGTTAACTGTGTTATTTTTGTCTGTATTATGTCAAATAATTTTGACTAAATCATTGGTATTATCAAGGTCATGTA
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Fig. 6.250

GAAGTCTTCTTAAATTCTGATTAAATAATCGACTTTCTTTACAACTACAGACAAATTGCAACAGAATCTGTATTTTCAG
GCAGTTCAGCCTGCATAGGGTCAATTTAATACCATATAAATGTTGGGAACAGAATTATCAGAAACTTCAGATATGTCAC
CATAAGCCAGATGAACAATATATGTCAGTACCATTGGGGGAGTATTTGCCAGCATAAACTGGATATGTTTCTTGAATGC
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CAGTTTTACTTGAAAGGCCTTCAGATTAAATACTTACAGAACCAGGAAGAAATGTATCTGAGTTACAGAACTTGAATGG
GGATTGTCAAACTGGAGAATGCTTGCCCCATCTGAAGGAGATAGCTGGCAATATGTAGGGAGGGGAACTCTAGGTATTT
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CTGCAAGCTCCGCCTCCCAGGTTCATGCCATTCTCCTGCCTCAGCCTCCGGAGTAGCTGGGACTACAGGCACCTGACAT
CACGCCCGGCTAATTTTTTGTGTTTTTAGTAGAGACGGGGTTTCACCGTGTTAGCCAGGATGGTCTCGTTCTCCTGACC
TCGTGATCCGCCCGCCTCGGCCTCCCAAAGTGCTGAGATTACAGGCGTGAGCCACCGCGCCCGACTGGGAACTGAAGGT
TTTAAACCTCCTAAACATTCAAAAGAAGCCCAAATCTCAACTTTAATGTGACATTTTCTGTGTGCTTTAAATGTTGGAT
CAAAAGGAAAAACAAAATTAAAAATATGTTGGACAAACTCAGGCCTGCTGTCCATATTTGATCCAAGGAGGACCAATTT
ACCCACTCTATCCTACACCTTTCAGGTCATAGCATTTTCCTTTGTTATCTTTCGGCATTAATGTCATATTCCAATTTTT
CCATATCTATGAGCAAGGCCCAAGGTCTGTAGCCAAGCCCAGGGTTTGCATGTGATATCTTTAGAAAAACCCCAGAGAA
GCTTTCTCTTGTTTTCTCACTATGCTGTAAGTATGAATAGCCCAAAGCAAAAAAGACAAGTACCCTGACTATCATGATC
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GACTGGTTATCTCTTAGACTGTATGACAGCAAAAATATAAGCAAGGGTTAAATTTCAAATGCATCCACTCTGCTCATAC
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TGTTCATGATGGCAAAATAATCACGTTAAATAAATTCTAGATACTGCACTCTTATTATTAATAATAGCAATGATATTGT
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CTTGCTGATTTGTCCCTAGTTTATCTATTATGAAACTTGGATGTTGTCAAACCACATTTTCATTCCCCAGTCTGTGTT
CACTATTTGATGTTCCCCACAAGTTACACTGCCTTGGTCTCAACTAAATTGGTTTTCTATATTACAAAGTCAATTTAAT
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ATAGATGTACAGCAGAGAAATTAAGCTGACTGACATTCTGTGCCCTGCAATCGTCATATCACACCTGGGGTGTGGGATT
CATTTCAGATACCACAGGTTAAGAGGAAATCAACAAGGAGCAATCAGTCCAGAAGGGTGTTGCCAGGATGGTGAAGCAT
CTGGGAATCATGATGTCTTCGGAACTGCCAAAAGAACCGAGACTGATGTGGCCTGGAAAAGGGAGAGAATAAAAACTCC
TGTTTAGATAATCTGAGGTTTTCATATTAAAAAGATTCAGGGCCAGGTGCAGTGGCTCACGCCTGTAATCGCAGCACTT
TGGGAGGCCAAGGCAGGTGTATCGCTGGAGTCCAGGAGTTCAGGACCAGCCTGGGCAACATGGTGAATCACTGTCTCTC
CTAAAAATTTTTAAAAAATTAGCTTGTGCAGTGGCCCATACCTGTAATTCCAGCTACTCAGGAGGCTGAGCTGGGAGGA
TTGCTGGAGCCCAGGCTGTGGAGGCTGCAATGAGCCGAGATCACACTACTGCACTTCTGCCTGGGTGACAGAGTGAGAC
CCTGCTTCTAAATAAATAAATAAGACTCAGGCTTGTTTTTGAATGACTATGATCAAAGAATAGCACTTTTAGAAAAGGTG
AAATTTAGTTTGATACAAAGAACTTTTCAATATTTAAATCTTCAAACTTTTCAATATTTAAATTTTCGATATTTAAATC
TGCTCAAAGATAAGCATTTAAATCTGCTCAAAGATAAGCATACCATGAAGCAGGGCTCCTCATGCCAGAACCAGATATT
TGTTTTTCCTTCAAGGCTGTTCATGCATTAGGGGAAGACTGAATTTCCCATGGCTTCTAATGAACCTTTTAACTATGAG

Fig. 6.251

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GATCTGTTACATGAAAATTTTTTTTTTCAAGATGGAGTTTCGCATTTGTTGCCCAGGCTGGAGTGCAGTGGTGCAACCT
CGGCTCACCACAACCTCCGCCTCCCAGGTTCAAGCAATTCTCCTGCCTCAGCCTCCTGAATA. TGGCATTAGAGGCAT
ACACCACTATGCACAGCTAATTTTGTATTTTCAGTAGAGATGGGGTTTCCCCATGTTGGTCAGGATGGTCTTGAACTCC
CAACCTCAGATGATCTGCCCGCCTCAGCCTCCCCAAGTGCTGGGATTATAGGTGTGAGCCACCGCGACCAGTGTTACAT
GAAAAATTCTTAGAATAGTGCTTGATGGCACCAAGTAAATGTTCAATAGATATTAGTTGTAATTGTTATCAGTAGTGAC
TATTCTATGCATTTGTAACTGATTATTTTTGGATGGGAATAAGTAAAGGTGTTTAAAATGTTAGCACCGGTTTCTGTT
.TCCATTATTGCCTTTACATTCTATGTTACCTAGATTTGGGATTAATTCATCAAATTTAATATGAAACTTTAAGTGACT
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GTATTCATTCATTCACTCACTTCCTAATTTATTTAACAAGCAGTGTACCAACCACTGCCCATGCCTGCACCTAGCTGCT
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CTATTTAATCATCATTCCAGGGCAATAGCTGTATACTATTCAGACAAACTGAAGAGTAAAGTCCCCTAGTTTAAAAACT
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TGACTAGCTGATTTATGGGATTTTATGTTCTTTAACTTCGAAGAAGACATACAATTCAGTCTCAACAGTTTTTACAATC
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TCTGACTGTAGCCATTGTGTGTAAGACTGCAAAGTGTAATATATGCAGTGTTTAAGAGTAGATTAACAAGAAAAGCTAA
TGTATGAACATAAGTAGCTGACCCTTAATAGAAGAGTGTTTAATTAATTTTGAATTTGCCAGTACACTCTCAAAACACA
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Fig. 6.252

GAGCTTGATTAAATAATGGCATTATACTGAGCTTATTCAAGTATTTGGATAACTTTTCTTTACTGAACTGAAACTGATA
GCCCAGGCAGGCACATCTCCAACACCTCTAATTAAAATCCACATTCATCCCTTGCTTCCTGAGAAAAGATGTTGCTCTC
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CAGTTGTCAAATATTTATAATTATTATTCTTGGGAAAGTAACTGAGAAGGGAGTTCAGGAAAGACTTGGACAGTTCTTT
GGAGCATCAGGACCTTACAATTTCAGCTATCTTGTTCAATAAGCAAATGTTGTTTTAGGTTCCTGGATTACAAGGTCAA
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AAGTCATGTGAGTTGGCAAAATAGATGAGAGGCTTCTAAGTAAAACTGAGGAATGAAGAATGTTTTTTGGAAGAGATTA
CGCTTAAAAGGAATTAGCCAGCCAGAGAAGAAAGAGAAAAATGTTCTTGCCAGAGTACCTATCATTTGGGAAGCCCCAG
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CAAAACAATAGAGCATCTCTGTGCTGAGTATTGGAATTTAAATTTTGAACTTCATTACTAAATAATGTGGAACTATTGG
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Fig. 6.253

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Fig. 6.254

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Fig. 6.255

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AGTTATGCAGAGTTGAAGATGAAGGGCATTTCAGGCTGGGAGAACAATCTCACAAAGGTGTGGATGTAGGAAAATACAT
GAACGTTTATAGAAAAAAAATTGAGTGTGGTCAGAATTTATTGAACTTGAGAGGCAATCATGTAGATAGGAGATAAAAC
AGGAAAAGGAGAATGAGACTATATTTTTAGGGTCATGAAAATTAAGCAAAGGAATATTTTCTAAACTAAGGTGAGCTAT
TAAAGATTTTGAAGAGAGGGAGTCCTATGATCATAGCTTTGCTTATGAAAGAATGAAGTGGCAGCAATGAAGACACACC
TTGGAATACCAGAAGACTAGAGTAAGGAGACTAATGAGGGGACTTATAGCAATCATTTGCTTTGCAGGTAATAAGGTGG
TAACTAATACAAGATGATGGATTACAATATTAAATTTTGTATTCTTTTAAGTTCTATGTTTCTGCAATGGCTAAATGCA
AACCTATCAGGAAAAGTAAAGCGTTTTGTTGTTATTGTTGTTGTTTTTGCTTTTCAAAAAGTGCAGGTAATTAGGGCCT
AACGTGGAATGGTCCATGCTAGGAATAAAGTAGATAGCGGCGAATGTTTGCTAGAGACATTGTGATGGACTGATCTGCC
CTACTATTACATCCTCAGCAATAAATATGAATGTTTAAGTGGTACAATTGCCAGAAATCAGCCAAAGTTTGGCATAATT
GTTAGAGATTTGTTGGCTGTGGATAGAACTACATAATTTTGGAATTGTAGAGGAGAACAGAGAAGAAACCAACAAAAAA
TAGTACTAGTACTCTGTCACTGGAAAGAAGTATGTTTAAGGCCACACAGTGAAAGTTAGCCATGAGCTTGAGTGCTCTA
TCATTTTTCATTATGTTGTTTGTTTATTAAAAAAAAAAAAGGTAACAGCTTTTTCTTTTTAACCAAACATTTCTTACTG
GAAGTTCAATAGGTGTACAGTTGTTTTGTTTGGCTAGACATGGATATTTGTGTGTTATTCCCTTTCAGTAGTTCTGAAA
ACCATTTTATCTTTAGATACATTATTTTCCCAGGAGCTTGGGTATTTTATTGAAGCTGTTTCAAATGCATTTAATGTCC
TTTGTAATGGATTTCTTTATCTCTTCCCCAATGCTCTTGGCTGGAGATGTTACTTTTTATTTGCCTTATCGGTGTGAGC
ACCTCATTGCTATAATCAATAGATATAGTACTTTAGCATTCTGTACATTTTAATATGATATATACCAAATATAATGTAT
AAATGAAAAGTTATAGATAATTTTTGCTTAAGTTTTCTTTTATAGAATTGTTAACAAAGGATATACAGCCAATATGT
```

Fig. 6.256

TAAAATAATACCTAGAAATTAAAAAGGAGTAAAGTAGAATAGTTTATCTGTTGTACTAAGACTTCATACACAATATTTC
GAAAAGTGGGAATATCTCTTGGGTGAAATACTTCATATATGTATTGTATGTACATACATGCAATATGTACACACAGAAT
TATATAAAATATATGTATATGTATACATCTATATGTACAGAACATATAATTTTATTAGCATTTCTTGTAATGGCATTTT
ATTATAAATCCAATGACCTCAATTATTCTATGGGTAAGTGCTAAGTATGTCAAGAGAGCTGGCATAGAAAATGGAAAAA
GCACTACACCCCTTAGCTTGCAAGTAGATGTGAATTTTCTGCCTTTGACTTTGTTGAATCTGTGATGAATCATATGTTT
ACTCTGATTAACATAAACATCTGGATGATCTAACTTTGGGGACACATTGCTTCATATGCACTGAATGCCTGAAAATTG
GTAGAATTTTAGATTCTTTTTCTTTATAAATGATACTACCCGAATTCCTGCAATACCTAGAGAGTTACAAGTGCTTAGC
TCTGACCTTTTATTCCATTCAATTGAAGTTGTCCACCTTTAGTTTATTACACATATGACTCTTAGTAGAGCAAACATCT
GGATTATTGTCAACAGTTCTCAAACACACCATTTGGATTTCACTATCAGACTACGAACGAACCCCCATGGAAAAAAAATT
CAGGCATACAGGCTACACCCAGATCCTGAATAGCCCCTGGTTTTCTGGTTACTATTTTCTCAGGCCAGATCCAAGAAGT
CCTCTTTGGGCTTGTCTCTGGGATTCTCTGATAAAATTGGCTTTAGATTGAGACTGACGTGAAGAGAGCTGGTCATT
GAAAGACAGAAACAGATGTGAATGAAATAATTCTCCTTTGAGACATAAAAAAATGTAAGATATACCAAGAAAGGGGAAT
TTGAGTTGTTCATTTTCTGACAACTGTGAAATTGTTGACCCGGACAAGAGAGTAGGGGAGATTAACAATGTGATCATGTT
CAGGGCCCACTGTGCCACTGAATTAAGACAATGTTGGCATTCTCCTCAGTCATCTTCCGAAAATAGAGATTTAGAGTTA
GGAAAGGAATAGCCTATCCCATTAACTACGTCACTCATCTCATGCAGGTGACCTCTCATTTATGTATTTCTCAGGATAT
AGACCAACGCTGGAGTATAGAAATATTATGAGACACTTCGTAATTTTAGGTCTTCTGGTACCACATTTAAAAGGTAAAA
AGAAATAGGTAAAATTTATTTCAAGGATATATTTTATTTAACTCACTGTATCAGATTATTTTAACTTGTAAACAATATA
TATAATTATTAATGAGATATTTAACATTCTTTCCTCATACTTTGTCTTTGAAATCCTGTATGTATTTTTTACTTACATC
ACATATTAATTTGGATGCCCAATTGTCCTCAATTATACTGGATCTTTATTGGACATTTAGATCTCATAAGATTTACAAT
TAGAAATATAGATTCACTTCCCAAGTTGTTCCAAATATAGTATTATTAAATCAACTATCAGTCTTTTAATTCATAAATA
TTAAGTAAGACTAAAAATTTAGTCCTTTAGTCATACTAACCCCCTTTCAAGTGCCCAACAGCCTCTTGTGGCCATTGGC
TGCCGTATTAGACAGTTCATATTTAGATTATCTGGGTTTGAACTCAGAGTCTGCCACTCACTGACTGTATTGGCAAGTT
GTTGCCAACCTGGGCAAGTTTTTTTCTCTTTTTATGCTTCAGTTTCCTCACCTGAGCACACCTTTTTTCCTTGAAGAAGCT
CTCCTGAAGAAAATCCCGTATTTTTCTTCAGGACATTGCTATTTTTCCACTTAAACTCAGAACCTTAGAGTGATCACTG
AATCCTCCTCTCCCTCATCAGCTTTATCTGCCAGAAGGTCAAGTCCTCTGAGATGTACACTGTAAAATCTCATGCTTTC
CTGTGACTTCATTTCCACTGCCACTACCCTCTCCTTTTGCATGGATAGCTGCAATGATCTCCTATTTAATTAGCCTCAG
TCTCCAGACTCTAGCCATCACTCCACATTATGCCCTCTGGATGCTGCTACTGAGTTCATTTTCTGAAAATGGTATCTTT
ATCATACTCTCCACAGTTCAAACTATTTCTTCACTGCCCGTCACTTCCAGCATAAATCCATGTCCTCTGTGAAACCTTT
TTTGACAACTTCCATGCCAGATCCAAATGACAACTCCCTTCCTTGTGCTGTCATGGCCCTCCACACAGACTTCTGCCAT
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TATGAGCTAGTGAATGACATTCCTTTGTCTGGCATTCCAGGCCTTCAGAATTAGCTCCACTTGACTCTTCTGGCTCCTT
CTCTTGCTTTTCATCCACACAGATCCTCTGGTATAGACAAACTGGTGTATTACTGTCTCACAAATATGTCCCCAATGTT
TGTGTCTCCATGCCTTGGCTCGCATCCTCTCCTCTTCTGAAGTGTCCCTGCATTCTTCTTGATCTGCCCAAATTCTACC
CTTTCTTCAAGGTCCAGTCCCTCCCTCTTCCATGATGAAGACTGCGGAGCATGATGCAATTCACTGTGTTTCAAATTCA
GGATTGGGTGGCCTGATGGAAGGAGGCCAGGCTTGGGGTCCAAGTGGATCTGTTCTTCTGTTTTCCGCCAAGTTGCCT
ACCCTCTCTGAGGTTCACATTCCATATCTTTGAAATCACAGTAACAAAACATAAGGTGCAGAACTATCTTGAGTATTAA
GAATAACGTGTGCACTGGCATTCAGTGAGGGTTCAGATGGAGCACGTGAACTCTGTTACTTTCAGTACAGTCATAGGAT
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TTTCTAAACTAGACTGCAAGTGTCTGGACTGAAACAAATGCATAATAGAATTCTGAGCAGATATGCACAGATATATATA
TGTGTGTGTGTATGTGTATATTTGTGTGTGTGCACATGTATTTGTAGATATCAAGGGAGAGAGGCACTTTCATGTGTAT
TGATTCATTCAGTCCCCACAAGAAAGCATCATTCTAGGAGGAAGTTGAAAATGTCCCTATTTTATCTATGAAGACTCTA
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TAACCATCTGTCAATCTTTATCTACATGCAATTGTACCCAGGTTCAGGGATTGTGGGAGGGAGAAAGGTTAGGTGGTTG
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TTTTTTTTTTTTGAGATGGAGTCTCACTCAGTCACCCAGGCTGGAGTGTAGTGGCGCGATCTCGGCTCACTGCAAGCT
CTGCCTCCCGGGTTCACGCCATTCTCCTGCCTCAGCTTCCTGAGTAGCTGGGACTACAGGCCCCCACCACTACGCCCGG
CTAATTTTTTTGTATTTTTAGTAGAGACGGGGTTTCACCGTGTTAGCCAGGATGGTCTCGATCTCCTGACCTCGTGATC
CGCCCACCTTGGCCTCCCAAAGTGCTGGGATTACAGGCGTGAGCCACCGCGCCCAGCCAACCATGGCATTTTTTGTTAG
GTCACTCAGCATTGGTATCAAGAATAACAAATGACACTTGAGTTTCTTTTTCCTGAAAAAGGGCAGGAAGAGTCTAATA
GCAAGTGCAATTGCCACAGGCAACAGTGTTATAACTGGAAAATCCTAAAATGTAGATAATTTTCCTCCAAATGCTTTCT
AACAAAAGATAGACAAGTTTAAATTTGGCTGATTTCATATCTACCATAATATAGTACCAGCAATGCCAAGATACAGAAA
ACTGCCTAAGAAAAACAGAGTTCAGAAGACTTCGGATAAATTATCATCTTAAGCTTAACCTTATACCTATAAAAAATTC
AGGGATATCAGTTATGACTTCTCATTGAGTAGTCTCATGTTAGACCAAAATAGTTTCCCATATTTTGGTGAAGGACCAG
AATAACTTACCAGTAAATGAAACAATCATTTTCTTTTTGCTTTATACTCATTCTGCATGTGATTTGTATAGGGGATCAA
GTCAAAGATTGCCAGTACAAAGTAACAACCTCTCATTGTATTGCTTAAGTTAATCATTAATATTTTCCATGGATCAATA
CCCTGTAGAAGCATGAGATGCAGCAGTGATCTTCAATTTCATGTGCTTGCCAAGTAAGAACAGCCATGGGCCAGATTGT
TGGGAGCCGTGCCATGAGCTACAGACCCTCAGCTCCCCTCTATAATCAGTTCTTCCCCCACTCCAGTGCTCCCAACTTG
CCCTAAGCAGAGCGTAATTGTGGATGTGTAACTACCGCAGAGGGGAGTATGCTTTTTATGTTGTTCAATTCTTCACTTT

Fig. 6.252

CTCTTCTGCAATTGAAAGTTGAGCTGTTAGATTTCCTGAAATGAAATGGTGACAAGAGAAAGTAACAGAAGTAGCCTTT
ATAAGTGCCTGACTTCTGGGTAACAGAAGTGACTTAAAAAAAATAAAAGTACAAAACGTTGCATTTAATGTAAAAGATG
ACTGCATTAGTACAACCCTTGAAAACATTCAAGTTCTTTTCAGAATAACCACCATTAAGGAGTTGACAAATATTTTATG
CTTTAAAGTATCCTATAAAAGTTCATCACAGAAATAAAAAGCCTCCTGTTCTTATCCTTCTCAACAAAGCCCTCTTGTA
TATAGATATTGAGTCTGGCTTACCTCTGTATCCTGTTGTGCTGATAAACCAAGAACCTTGCAAAAATAGATGCACGATA
ATTGATATTTAATAAGTGAGCGAATCCATTAATGAAAGAGTGCTATTTCAGAGAGCAGTGTTCATTTAGATAGATGGT
CCCAGCGTCATTAAGTAAAAGGATAGAGAATACTTGCAATACATTTGGCGTAGTGTTAAGAATTTTTTAAAATGTTTTA
AATGTGATCTTTTTTATGGGCTGAATTGTGTCCTCTCAAAATGTTGAAGTCCTAACCCCTGGTACCTCAGAATGTAACC
ATGGAGATAAGGTCTTTAAAGAGGTGATTAAATTAAAATGAGGTCTTTAGAGTGGCCCCTAATCCAATCTGAATGATAC
CTTTATAAAGGGAGGGAATTTGGACTCACAAAGAGACATTAGACACATGCACACAGAGACAACCATGTAAACACATGGT
AAGAAGGCAGCCATCTGAAAGCCAAGGAGACAGGTCTCAGAGAAACTAAACCTGTTGACGCCTTTGATCTTGGACTTCC
AAACTGTAGAACACTGACAAAACGTATTACTGTCGTGTAAGCCACCCAATCTGTGGTATTTGTTATGGAAGCCCTGGA
AAATTAATACAGTCTTTTTTATTTGCTACCTTTGAAAGCCCTCTTGTCAAATGACGGTAAAGAATAGAAGACAATAAAG
TGTCAGTAATAAAGGGTGACGAGAAGAAACAACTAGAGATGTGAGAGGACCAATTTAGGAAATGGAAAGGAGAGGAAAA
AATCAACAAAGTTGAGAATGCTAATTACAAAGTGCCCACAGAGGAGGGCAAAACCAGCAAACCATTCCCTTTGTCAAAA
TCTTGAAAGATGGCAATTGAGGAACCAAGAATCTTAAAAAGGGGGGAAGGTGGATGAGAAATCTGCATATGAAGGGGAG
TCCTCCATAGGTTCTCCTCTCACCCTGCTCAGCTAGGAGACCAGTGCCACTTTATCCTCATTAGAATGGAAATTTATGC
TATAAAAATAAATAAGAGAAGTCTTAACCTGGGGATACTAGGCACATTGGAAGGTGAGAGAGTAAATGAAAGTCTATCA
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TTGCCAGGTTCAGGGGCTCACGCCTGCAATCCTAGCACTTTGGGAGGCTGAGGCAGGCAGATTGCTTGAGGCCAGGAGT
TCAAGACCAGTATGGCCAACATAATGAAACCCCATCTCTACTAAAAATATAAAAATTAGCCAGTCATGGTGGTGCATGC
CTGTAATCTCAGCTACATGGGAAGCTGAGGCACAAGAATCACTTGAACCTGGGAGGCAGAGGTTGTGATGAGATCGTGC
CACTGCACTCCAGCCTGGGTGATGGAGTGAGACTGTCTCAAGAAAAAAACAAAACAAACAAACAAACAAAACACTTGAA
CTCGTGGAGATAGAGAGTAGGATGCCATTTATCAAAGAGTGGGAAGGGTAGTGGGGAGGAAGAATAAGATCTAGTATTT
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TTGTAACACAAAGAAATGATAAATGCTTGAGTGATGATGGCTCCTCCATTTATCCTGATGTGATTATTACACACTGTATGC
CTGTATTAAAATATTTCATGTGACCCATAAATTCATATACTTACTATGTACTCATAAGAATTAAAATTAAAAATAAAAA
TAACCAAGAAGAATTCTTCTTCAGAAAATGGAATTACCACCCAGAGAAAAGACCTATAGTACTGGCATTAGATGGTTG
CCCAATAAAGAAGCCAAGGGTCTTTATTCCTCAATTACCCCATGTATTAGGCCATTATTGCGTTGCTATAAAGAAATAC
CTGAGAGACTGGGTACTTTATAAGGAAACATGTTTAATTGGTTCACAGTTCTGCAGGATGTACAAGCATGGTGCTGGC
ATCTGCTCAGCTTCTGGGGAGGCCTCAGGGAACTTGTAGTTGTTTCAGAAGACAAAAGGGGGAGTAGGTGTCTCAAATGG
TGAGAGCAGGAGTAAGGGGGTTGTGGGAGTTAGCACACACTTTTAAACAACTAGATCTCAGAGAGCTCATTCACTCACT
CACCAAGCCATAAGGGATCTGCCCACATGACCCAGACACCACGCACCAGGGCCCATCTCCAACACTGGGGATTACATCC
CAATATGAGATTTGGAGGGGACACCCAAACTGTATCATTCCGCCCTTGGCCCCCTAAATCTCATGTCCTTCTTACATGG
CAAAATGGAATCATCTCTTCTCAACAGTCTCCCAGAGTCATAACTCATTCCAGCATTAACTCAAAAGTCCCAAGTCCAA
ATACAAAATCTTGTCTGGAAATGAGTTCCTCCCACTTATGAGCCTGTAAAATTAAAACAAGTTATTTACTTCCAAGCCA
AGGTACAATAGAGGTACAGGCATTGGGTAAACATTCCCATTCTTTTTTTTTTTTTTTGAAATGGAGTCTCGCTTAGTCA
CCCAGGCTGGAGTGCAATGGTGCAATCTTGGCTCACTGCAACCTCTGCCTCCTGGGTTCAAGCGATTCTCCTGTCTCAG
CCTCCTGAGTAGCTGGGATTACAGGCACCCGCTGTCATGCCCAGCTAATTTTTGTATTTTTGTAGAGATGGGGATTTTC
ATCACATTGGCCAGGCTTGTCTTGAACTCCTGACCTCAGGTGATTCACCTGTCTCAGCCGCCCAAAGTGCTGGGATTAC
AGGCATGAGCCATCATGCCTGGCCAACATTTCCATTCTAAAAGGAAGAAATCAGCCAAAAGAAATGACTTCAGGCCCCA
CCCAAGTCTGAAACACAACAGGACAGTCATTAAATCTTAAAGTTCTAAAATAATCTCCTTTGACTCCATGTCCCACATC
CAGGTCACACTGATGTAAGGGTAGACTCCAAGGCCTTGGGCAGCTCTGCCTCTACAGCTTTTGCAAGGTGAAGCCTCCA
TGACTGCTCTCATGGGTTGGAGTTGAGTGCCTACGGCTTTCCCAGGTATAGGATGCAAGCTGCCAGTGGATCTACATTC
TGGGACCCACAGAACAATGGCTTCCTTCCCACAGTTCCACTAGGCAATGCCCCAGGGGGGATTCTTTGTGGGCAGGGCT
TCAATCCCATATTCCCTTCTGCACTACCTTAGTAGAGGTTTTCTGTGAGGACTCTGTCCCTGTAGCATGCTTCTGCCTG
GACACCCAGGCTTTCTGATACATCCTCTGAAATCTAGGCAGAGACTAGGATCACTCTTTCCATTCTTTCTATCTGAAGG
CTTAATACCACAAGGAAGCCACAAAGGCTTACAGCTTACATGCTCCAGAGTGGCAACCTGAGCCATACCTGAGGCTCTT
TGAGCCACGCCTGGACTGGATGGACCATGATGCAAGGAGCAGCCTCCTGAGGTAGACAGTGTAGTGGTGCCCTGTGCTT
ATCCCCTAAACCATTTAGTCCTCCTGTACCACTGGGACTGTGGTGGGAGGAGCTGCCTAGAAGATCTCTGAAATGCTGT
AAAGGCCTTTTTCCCATTATTTTGGCTATTAGCACTTGTTCTCTTTTAGTTATGCAAATTTCCCTAGCAAATGATTGCT
TGCAGCCTGCTTGAATTCCTCCCCAAAAATGGCCTTTTATTTTCTATCACATAGCTAGGTTGCAAATTTTCCACACTTG
TAAGCTCTGCTTCCCTTTTAAGTAAAAATTCCAAGTTTAGGTTATTTCTTTGCTTCTGCATCTGAGCATAGGTTATTAG
AAGCAGCCAGGTCACATCTCAATTGCTTCAGTGCTTAGAAATGTCTTCTGCCAGATACCCTAAGTCATTAACTCTTAAG
TTCAAACTTTCACAGGTCCCTGGGCATGAATATAATGTAGCCAAGTTCTTTGCTAAGGTATAACATGGGTGACCTTTGC
TCCAGTTCTCAATAAGTTCCTCACTTACACCTAATACTTTGTCAGCACGGACTTTACTTTGCAGATCACTATCAGCATT

Fig. 6.258

TTGGTCATGACCATTTAATCAGTCTCTACCAAGTTTCAAACTTTTCCTCATCTCCCTGCCTTCTCCTGAGCCCTCCCAA
TTCTTCCAACCTCTGTCTCTTACCCAGTTCCAATGTCATTTTGACATTTTCAGGTATCTTTATAGCAATGCCCCACTCC
TTGGTACCAGTCTTCTATATTAGGCTGTTGTGGCATTGCAATAAAGAAATATCTGAGACTGGGTAATTTATGAAGAAAT
AGAAAATTAACTGGCTTGCAGTTCTGCAGGCTGTACAAGCATAATGCCAGCATTTGCTAAGCTTCTGGAGAGGCCTCAG
GGAGCTTTCAATCATGGTGGAAGCTGAAGAGAGAGCAGCTATCTCACGTGGCAATAGGGGGAGCAAGAGAGAGAGCGGG
AAGGAAGGTGTTACACACCTTTAAACAAAGAGATTTTATGAGAACTTGCTCACAAAGCTATAAGGGAACCACTCCCATG
ACTCAGACACCTCCCACCAGGCCCCACCTCCAACACTGAGGATTACATCTCAACATGAGAATTGGAGGGGGACATTCAAG
GAACACCACCCTATAATGAACCCCAAGACTCTACAAGTCCAGTTAAATCATTCAAGGCTCCCAGTCAGTATTTTAGTGT
CTCCCTGTTGAATATGAATAGATAGCCAAGTCTCCACAAAAGAGTTTAAGAAACAAATAGACGTTTCAGAAAAAATAAACTT
CGTAGCCCTAGACAGATGACAAGTTATCACATTAATGAAACAAAAGGATACTATACAAAATTATCAAGTAAGAACAAAA
GAGAGTTAGAATTTAAAACAATGAAAGAAGATGCTAATAAATAAAATCAGAGATGAAAAGGGAGATGTTGCTGGGCACA
ATGGCTCACACCTGTAATCCCAGCACTTTGGGAGGCCAAGGCAGGCAGATCATGAGGTCAGGAGTTTGAGACCAGCCTG
ACCAATACGGTGAAATCCCATCTCTACTAAAAATACAAGAATTAGCCAGCGTGGTGGCACGTGCCTGTAATCCCAGCTA
CTCAGAAGGCTGAGGCAGGAGAATTGCTTGAACCTGGGAGGCGGAGGTTGCAGTGAGCCAAGATTGTGCCACTGCACTC
CAGCCTGGATGACAGAGCAAGACTCTATGGGGGGGGGGGGGGGGAAGGAAAGATAAGGAAAGATAAGGGAGACGTTACA
AGCAATACCACAGAAATTCAAAGGATCATTAGATGCTACTAGGAGCAACTATATGCAAATAAATTGGAAAACCTAAAAA
AATGATAAATTCCTAGACAAATACAACCTACCAAGATTTAACCATGACGAAATCCAAAACCTGAACAGACCAATACCAT
CAAGATTGAAGCCATAATGAAAAGTCTCCCAGTAAAGAAAAGCCCAGGATCTGATGGCTTTACTGCTTAATTTTGCCAA
ACATTTAAAGAAGTAATATCAATCCTACTCAAACTATTCTGAAAAATAGAGGAGGAGGGAGTACTTCCACACTTATTGC
ACAAGGCCAGTATTACCCTCATACCGAAACAGACCAAAGGCACATTGCAAACAGAAACTACAGGCCAATACTCCCAAT
GAACATTTATGCAGAAATCCTCAACAAAATACTAGCAAGCCAAATTCAGCAACACATTAAAAAGATTAGTCATCGTGAC
CAAGTGGGATTTTTCCCAAGGAAGCAAGGATGGTTCAACATAATGTAAATCAGTCAGTGTGATACATCATATCAATGAA
GGACAAAAACATATGATCATTTCAATTTATGCTGGGAAAGCATTTGATAAAATTCAACATCCCTTCATGATAAAAACTC
AAAAAACTGAGGATAGAAGGAACATACCTGAACACAAGGAAAGCCATATATGACAGACCCACAGCTAGTATCATATCGA
ATGGGGAAAAATTGAAAGTTTTTCCTCCAAGATCTGTAACATGACAGGGATGTCCATTACCCCTTGAGCAATCAGACAG
GAGAATGAAAGGATATTCAAATTGGAAAGGAAGAAGTCAAGTTATTCTTGTTTGCAGATGATATTATTTTATATTTGGA
AAAACCTAAAGACTTCACCAAAACACTACTAGAACTGGTAAACAGATTCAATAAAGTTGCAAGACACAAATTCAACATA
TAAAAATCAGTAGCATTTCTATATGCCAATGGTGAACAATCTGAAAAAGAAATCAAGAATGTAATCCCATTTATAATAG
CTACAAATAAAATACCTAGGAATTAACTTACTGAAAGAAGTGAAAGAGTTCTACAATGAAAACTATAAAACACGGATGA
AAGAAATTAAAGAGAACACAAAAAATGGAAAGATATTTCATGTTCATGGATTGGAAGAATCAATATTGTTAAAATGTAC
ATAGTACCCAAAGCAATCTTCAGATTCAATGCAATCTCTATCAAAATACCAATGACATTCTTCACAGATATAGAAAAAT
ATCTTAAAATTCATGTATAACCACAAAAGACCCAGAATACTCAAAGCTATACTGAGCAAAAAGAACAAAACTGGAGGAA
TCATATTACCTGACTTCAAATAACACTACAGAGCTATAGTAACCAAAACAGGATGGTTCTGGCATGAAAACCCAGGAAC
AAATTCATACATGTACAGTGAACTCATTTTTGACAAAGGTGCCAAGAACATACATTGAGAAAGGACAGTCTCTTCAATA
AATGGTTCTGGGAAAACTGGATATCTATATGCAGAAGAATGAAACTAGACTTCTATCTCTTGCCGTATACAAAAATCAA
ATCAAAATGGAATAAAGACTTAAATCTAAGACTTCAAACTATAAAACTATTGAAAGAAAACATTGGGGAAACTCTCCAG
GACATTGGCCTGAGCAAAGATTTTTTTTTTTTTTTAGACAGAGTCTCACTCTGTTGCCCAGACTGGAGGCTGAAGTG
CAGTGGTGTGATCTCAGCTCACAGCAGCCTCTGCCTCCTGGGTTCAAGTGATTCTTGTGCCTCAGCCTCCTGAGTAGCT
GGGACTATAGGTGCACACCACCATGCTGGGCTAATTTTTGTATTTTTAGTTGAGACAGGGTTTTGCCATGTTGGCCAGG
TTGGTCTCCAACTCCTGACCTCAAGTGATATACCTGCCTAGGCCTCCCAAAGTGCATGGATTGCAGGTGTGAGCCACTG
CGCCTGGCCTGAGCAAAGATTTTTTGAGTAATACCCCACAAGCACAAGCAACCAAAGCGAAAATAGACAAATGGGATCA
CATCAAGTTAAAAATACCTGCGTAGCAAAGGAAACAGTCAACAAGGTGAGGAGACAACCCACAGAATGGGAGAAAATAT
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ATCCAATTAAAAATGGGCAAAACACCTGAATAGACATTTCTGAAAAGAAGACATACAAATGGCAAACAGTCATATGAAA
AGGTGTGCAATGTCATTGGTCATCAGGGAAATGCAAATCAAAACTGCAGTTAAATATTATCTCACCCCAGTTAAAATGG
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AGCAATCCCACTTCTAGGTAGGTATACACCCAAAAGAAAGGAAATCAATATATCTGCACTTCCATATTTATTGTAGCAC
TATTCACAGTAGCCAAGATTTGGAAGCAACCTAAGTTTCCCTCAATAGACGAATGAATAAAGAAAATGTGGTACATATA
CACAATGGAGTACTGTTCGGCCATAGAAAGAAATGAGATCCTGTCATTTGCAACAACATGGATGTAACTGGAGGATGTT
ATGTTAAATGAAATTATCCAGGCACAGAAAGATCATCTTCACGTATTCTCACTATTTAAGGGAGCTGGAAATTAAAAGA
ATTGAGATCCTGGAGATAGAGATTAGAGTGATGGTTACCAGAGGCTAGGAGGTGTAGTGGATGGGAGGAAGTAAGAAGT
GGGGATGTTTAATGGGTACAAAAATATAATTAGATAGAACAAATAAGATCTGGTATTTGATAGCTCAACAGGGTGACTA
CAGTCAACAGTAGTTTATCATACATCTCAAAATAACTAAGAGAATATAATTGAATTGTTTGTAACTTAAAGGGTAAATG
TTTGCAGTAATAGATTCCCTGTTTACCCTGATATTATTATGCATTGCTTGCCTGTATGAAAATATCTCGTGGGCCCCAT
AAATCAATCAATCTATCTATCTATCTGTCTGTCTGTCTGTCTGTCTGTCTGTCTCTACCTACCTACCTATTATGTAC
CAACAGAAATTTAAAATTTAAAAAATTATGTTTTGGAGACGTCTAGAAATAAAACATAATTATGGCACATCATTTGGTTTA
AGAAGGTAATATATAACTCCAGAAAAAACAAAAAGTTGTTCCATAATAAAACATAATTATGGCACATCATTTGGTTTA
TCAGTAAACGTTACTTTGCAGGGATGATACAATAGAACACTGAATGTTTATTTAGCCAATAATTGTTATATAACTGTAA

Fig. 6.259

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TGGGAGGATAGAAAGAGGGAAGTAGAGAGGTGAAATGJTGAAAGAGTGTGAAATCTTCATCATTCATAATAGAAAGTCA
GTTAGAAATATGGAATTTAAAAAAAAATAGTTTGAGAGCCTTATGAAGAGGGTGGCACAAAAGACTCCTGCAACTTGTT
CTAAGCCCTGTGCATACAATTTGGGGTGTGTGGGTTAGTTTAATAATTATTTTGTAAATATAAAAGGAACCAAAGCAAC
CAAATTTCTGGATATAAATGTCCTGTGTTGATTATCATGATAATAGGTAGACATCTTGGATTGGAATTTTTTAACATTT
GAATGGAATACCCCATTTCCCCTTTAATCTTTGATAGGTTTAACATGGTCACTAACTGTAGAGTAGGGACTTGTCGTGC
TCTTTATTGGCAGAAGTGTGACTGGCTACCCCTGACCGTTCGGGAGAGGAGACATGTACACCTGGCCAATCAGACCAGC
AGATTCACTGCTGGTGATGAGTGAAGTTGCAGATGTCAGGGACAGCTTGTGCTGTTCAGGTTAAATTATCTCACTTCAG
CTCTCACTCTAAAGATGATTTATAACTAGTATTCTTCACTTTCAGTCTTTTCACTAGCAAGCAGGATACTTATGGAAAA
TAATTAGGATGAAAAGAGGTCACTCTATCACAATAGTGAGTTTTTGAATTCACACTTTAAACGAGATGATCCTCCTTAG
GAAAAGCAAAATAAAACGAATTTCCATGCCATTGTTGATTCTACTTGAGTTAAATACTAACATAAAAAGTTGGTTTTCT
GTGCATAGCCTGGAGAATCTTGTCTCTCAGAATTATCATGCCCCAGGCCATTCCTTGCAGATTTTGAATTCCCTTTGTC
TCAAAAACATCCAGAAAATCTTTTTGACCTTCGGGTGGTTGGCAGAACATGTGTTGTGAGGGCACCTCAGTTTAGAAGA
AAAGCCCTGAACTTAGTTCAAAAGAAGGGCTTTCCAGCATGCTATTCATTCAGCTTAGGACTGCCAGGCTTCCTGAAAG
ACAGGTGTGTCAGAAACAAAGTAAATAAACAGAAACTCACAATGTCCAGACCTTTCTCTCCAGAGCAGCACAGTTCCCT
TGCTGAGCGGCAACAGCCAAGGCTTTAAGCTGTTCCCTCTCCCTTTTCTGTTAATTGAGAAAACCAATTGCTGCAAGAG
CAACATCAGCAGAGCTAAATGTAAATGGAGTTTAATAGAAAGAGACAAATAACCCACTGAGAACCCCTAACATTTCAGT
GTAATACCCCAGAGTTCCACACCATAAACTTGGTATTCCAGTATAAAGCTCATCTCTCAATTGCACGCCACCCTCCCCC
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CCCTACAAAGGACATGAACTCATCATTTTTTATGGCCGCATAGTATTCCATGGTGCATATGTGCCACATTTTCTTAATC
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```

Fig. 6.26C

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Fig. 6.26j

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Fig. 6.262

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AAAATGAAATCAAGAAAAAAAAACCCAGTATAAGCATATTACTTAGAATTATAGAGTTAATTACCAGCAGAAACAAAA
TGACTTGAGAAAGACATTCACAATGAATATATATAGTATAGTTTTGCAGACTGTTTTTTGAATAAGGTTTTCAGTATA
ATTTTAAAATTATATGTGCATAATTTCTCTATTTATCCTGATGGGTGAGTTAAATTGCCCTGACAACCTCTAAATCTTG
TAACAACAAAGATGATAGTGATTGTTGTTTCCCATACTACATATCATCGACAGGGAGCTCTGCTGCACAGCCTCCTCAC
TCTGGTACAGAAACTGACAGAGTTACCATCATGGACTTTGTTGATCCTTCTGGCAGAAGGAGAAAGGGGAAGAGATGGA
GAGTCTCACACCAGCGAATCAACGCTTGGCTGCTGATTGGCCAGAATTAGTCACGTGGCTCCATTCAACTACAAGAGGG
TTATTAAGTGCATCTACATGTGTCCTGAGGGTGGAGAGCTACAAAGAGAGAGCCAGTATCGCTAATGGTTGCCTCAATT
TCTCTCATAATAATTGTTAAGTAAAAATAAAATAGGGCTTGGCGCAGTGGCTCACGCCTATAATCCTAGCACTTTGGGA
GGCCCAGGTGGGAGGATTGCTCGAGTCCAGGAGTTCATGACCAGCCTGGGTAACTTGGCAAGACACCGTCTCCTCAAAA
AATAAAAAAAACTAAAAAAATAAAAAAATAGCCAGGTGTGGTGGCATGGGCCTGTAGTCTCAGCTACTTGGAAAGCTGA
GGCAGGAGGATTGCTTGAGTCCAGGAAGTTGAGACTGCAGTGAGCCGTGTTTGCACCACTCGCTCTTCAGTTTGGGTGA
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Fig. 6.263

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CAGAGGGAGACCCTATCTCAAAACAAATACAAAAACAAAAACAAAAATGGTTTACTTCTAGCTAAAAATTTAATTAAAA
ATTAAAAATTCCTATGGATAATCAGCATGTAAATATTTGTGTGATAAATATGTAGGTATATA CAAAATATGTATATA
AACAAATATACATATGGATTATATATGTATAATTTATATTTTTATATTTGGTTTTTTCATATGTGTATTATTGAGTCCT
TAAACAAATGTTTTTTGAATAGCTTCTATTTACCAGGCATTGTTCCAGGTTCTGGGTATTCAATAGTGAACAAAACAGA
CACCCCTATTTCTGTGTAGCTTATATGGAGTGGTAGGGACAGAGAGACTCAGTAAGCATAGTAAATAAATACATTGTGT
AGCGTCTTAGAAGGCTCTCAGTCGTAGGGAACAAATAGTGCAATAGCAGGGGAGTTCAATACAGAGGCTGAGAATATGG
TTAGAATTTGCAATAGGGAAATCAGGTTAGAAAGGGCCCTTTGAGGCAAGGATTGAAGGGGAGTGGGGATAGATTCGTG
GGCAAATGGGCAGGGAATGCAAATATCCTGAGACTGGAGCAGGCTTGCGATTTTTCCAGGAATGGCTGGAAGGCTAGTG
AGGCTGCAGTGAGGGGACATAAGTGAGATCGTGAATGGGGACATAATAGGGGGCATCATGTGAATATATGGGATGTAGA
GACCGTGCTAAAGACTGCCATACTCTCAGTTAATTGATTGAACTCTGAGTGAAACACTTGAATTTACACCGGGAAATAA
ATTGCCCAGCATCTAGAAGATGATTAATATTTCATGTAAACCAACTTGGTCACACTATGAAGATGGAAATGTAAAACTT
GTTACTTTTCTATCCATTATTAATTGCTTAAAGCTCTTCCATGCTTTAGATACCGCACTAAGCATTTATGTTTCAGCAA
AGTATTGGCATAAAATAAAGCTTACAGGATGGTGTGAAGGCAATATTTTTTGAGTGTCTACAGCATGCCACAAAATGAC
TTCAATGCTTAGAATAATTGAATTACATAAATATAATGTCCCCA  TTATAGGCGAAAATACTGAGGCTCAACAGACAT
AAAAATGGCTTGAGTTACCAGGCTACAGTAGAACTAGGATTTCAGTCCAGGTCAGTCTGACCCCAAAACCTTTTCTTTTT
TCATTGTTTGGATTTAGCACCCCCAAAAATGTCTTGTAATGCAGATTTCCTGGGTCCAGGGAATTGTTTAAGTTTTTGT
TCTAAAAACTTTCTTATGAAGCAGATTACATATGGAAAATGACCCTATGGACTATGAGAATTGGCCTTTAGAGAATGTC
ACTGCCAGCCCTGTTGGTATCTATAAGAACCAATAAGCATTATTTACAGAGAGTGATATACACAGTGATAATTAAGAAA
TAACTATATTTATGATGACAATTTTGCCAAATTTACTAGAAAAAAATAAAATATCCTCTGATTTGAGATTTATTGTTCT
TTTCTTTTGCAGGTAAGTTTGGAAATTTTTTTTGCTTGTGGAATAGAGGTATGGTATCACTTTTTTTCTGAATTTTATTT
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GAAAGTAACATTGTGGGTTTGCGTTATTTGATTCAGAAATTTTACATATGTGATTCCATATGGTAAATCTATGCAGTGT
TTGTATTTGGGTTTGTGTTTAAATTATTTCTGTGTCAAGTATAAGGAAGAGGATTGTAGTGCGGGGTCTGTGTCAGGGA
AATGGGGGACTGTAAAGAACAGAGGAAAGAATATGCATGTAATGAGAGGTTGAAAGGCTGAGAAGGGCAGCTCCAGGCA
TGTAATCTACAATGGCCATGAATGCATGTCATGAGGAAAAGATGCCTTTTCAGGAACAAAATAATCAACTGAAAACTGT
TTTCATTCACTTTTCTAAGCCCCTAGTTTTTTGTTTTCTCCTATTATTGTGCAAAATTTCTGGCCAGACTATTGCTGAC
ATGTACCTTTGAATAAAAAAAATGGCTGTAACTATTCTGAAACATGTTAAGTTAGAAAGCAATAAAATAGATCCACTTC
CTGGAAAAAAGCTCTTATTAACTATATGTCAACTATTTCAGATAAGGAGTAGGAAGTTTCAGTAAAGCTTCTAGATACA
CAAAAGAGAGGATAAAATATTAGAGGGACTTAGAAAGACRTGATATTTGCACTGAGGAAAGAAGGTAGCCTGTAAGTTG
ACCACATAGGACCACGGCTTTATTATTGTGCACCGTGCTTTCATTAGTATATAAAAGAGGTGATTGTGTGAGATTTAAA
TCAAGCATGAACTTAAACTAAATTCCATATTAATTATAAACGGTATGTTTACTTTATTCAAAAAAAGATATATGTTTCT
CAAAATAATTGCAAATATTATTATTAATTATRTAAAAATAAAGAGGAATGAGGATACTAATCTACTCACCTAAGTTCTA
GCTAAGAGATGAGGTAGCAAAGGAGGAAGAAGTTCTTCAGTTCCTGGTCTTGATGGGTAAAGCGATTTGATCTCATTAT
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AATCTAATTATCAGAGCTCTAGCTCCAGGGTCAGCAAACTTTTTCTGTGACAAGCCAGATAGTAAATATTTTAGCCTTT
TGGGGCCACAAGAACTCTGTAGTAACTACTCAACTCTGCTGTTGCAGAGCAAATGCAGTTATCTGTAGACAATCTGTAA
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ACTTCTCAAAAGCTTTGATGAATTTAAACACTAGAGGGCACAAAAGGACATTTTTAAGTCAAGGACCTCTTGTAAACAG
AACACTACAATTATGAGAAACTGCCTGAAGTAGAGCGATGTGACCTCAATGTGCATAATTTGTTTGAAATGTCAGCCTG
TAGCTCTTGTGACCTCACTGACCACAAAAACTTGAAGTGGTTGACCACCTTCATGTTGCTGTAATAATAGCCACATGTC
TTTCTTCTTTTTTTCCTACTTCTTTCTCACTTCTTTCTCTCTCTYTTTTTTTTTTTTTTTTGATTGCTTTTGGCCCTTT
CATAAAATATGGCCAAAATTGAGGATTTATATGATGGGATTTTTTTTCATTTCCTAATAGTTTTTTTAATTGCTTTAGA
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ATAATAAAATAATGCATGTTATTTATGCATATGATTTATGCATATAATAATGTATATTGACAGATGTCATCTGTTACAT
TGATTTATGAACTAATTTGAAAGTTTTATTTCAGAAAAATGCTTAAAAATCAGCAGAAGCTTGAAGCCTCAAAATTACT
CCTACCCTTTAACACATGGACACAAAGATGGTAACAGTAGACACTGAGGATTCCAAAAGCCAGGAGGGGAAAGGGGAAT
AAGGGAGGAAACTCTACCTACTGGGTACAATGCTCACTACCTGGGTGATGGGATCAGCACTGCCCCAAACCTCAGCATC
ACGCAATATACTCATGTAACAAACCTGTACATGTAGCCCCTAAATCTAAAATAAACATTGATATTTTTAAAATCCTACC
TCTTAATTTAGTAATTCTGCTTGTAGATATTTGGATTGTGGAAGTAATTTGGGATATTAAAATATTTATATTCAAAGAT
ATTCACCRTAAAAATACCTATAATATTGAGAACCTAGGTATATTSTATGTATTTCACATGAACTAATGTTTAAATAAAT
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GCTTATAGTAAACCATGAATGCAAAGCAGAATCCAAAAATTATGTACATCRAATCTCAATTGAATGTTAAAATTAGATG
CGTAAATATATGTATATACATATAAAAATAAAGAAAATGAGCCAGATTTACAAGTTGCCTCTCGGTATAGATGATTTTT
AAAAAAATTTAATTTTCTGAATTTTTCCAAATTTCTATCATGAATATTCATCATTCATATGGTCATAAAAAATAAGTATT
ATTCAAACAGAAAACTATTTTTGCATTCTGACAGCACTTTTTTAGGTGTGATTCTTAGATCAGTGGTGCTAAAAGTCAG
AAGGTTATAAGGAAATTATTTTATGTATCAGGCAATCAATTTAAATGAATTATTGATTGTTACGCAGACACATCATTAGA
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Fig. 6.26ç

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GTCACTTTTTTAGCCCTGGGATGGGATGATTTGTCTCAGGTTTGGGTGTTGTGGCTTAGTTGCCGTGCACTAGGCAGTG
AAGACCAGGAAGGAACAAGGGCCTGCTGGAGAGGCCTGCAGTGCTGATATTGAACATGGACTTCCCTGTAAGCTCCCTG
GACTGTGGTAAAGCAGGACACATCACACACACACACACACACACACAAACACACACACGTGGACACACACACTCATGCT
CCTCTATCCTTTTTCTCTTCTAAATCTGCTTTCTGCCCTTGCCTTTCCCTRTAAATGGATGTGTGTGGAGAGGAGCTGG
AGAGACAGCAGGGGTCTGAAGGATGCTAACTTGGACTTTTTCCTGAGCAAAGAATTCCTTAAGAATGACCTGGTATAAA
AATTTTGAAGAAATCTCATAAATAATATGGGCAAAAATGTGACAGCAATGTGGCACAATGCAAAAGCTTGTCACTAGTT
TCAGCCCAGCTTTTCCTCGTGATTGTTTTGAGTAGCATGAATTATCCTCTAAAATGTTGGTTTTTATTCCTTTTTCATT
TTGTACAAACATACCTAAAAGGGTGAACATAGTATAAGCAAAGACACAACGTTGGAACTATAAGGAATAATTTTTTCTT
AATTCACTATGTATTTTTCTAACACTTTTCTTGGAGCTTTTAATCCTTAGAACCTAGGAGATGGAGGACTATACACTCG
GGTTTACTTTTCTTTCATTTTCTTCTTCTGAATTCTTTTTTTTAACTTTTAAGTTCAGGGGTACATGTGCAGGTTTGTT
ATATAGGTAAACTTGGGTCACAGGGGCTTCTTGTACACATTATTTTGTCACCCAAGTATTAAGCCTACTACCTATTAGT
TTTCCTGATCCTCTCCCTCCTCCCACCTTCTACCCTCTGGTAGGCCCTAGTGTGTGTTGTTCCCCTCTTTGTGTCCATG
TGTTCACATCAATTAGCTCCCACTAATAAGTGAGAACATATGGTATTTGGTTTTCTGTTCTTGCATTAGTTTGCTAAAG
ATAATGGCCTCCAGCTGTATCCATGTTCCTGCAAAGGACATGATCTCATTCTTTTTATGGCTGCAGAGTATTCCATGG
TGTGTATGCACCACATTTTTTTTTATCCAGTCTATCATTGATGGGYGTTTAGGTTAATTCTATGTCTTTGCTATTGTGA
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TGGGATTGCTGGGTCAAATGGTATTTCTTTTTTAGGTCTTTGAGGAATTGCCACACTGTCTTCTACAATTGTTGAACTA
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GCCCGCCACCACGCCCAGCTGATTTTTTGTATTTTTTAGTAGAGACAGGGTTTCACTGTGTTAGCCATGATGGTCTCGAT
CTCCTGACCTCGTGATCTGCCTGCCTCAGCCTCCCAAAGTGCTGGGATTACAGGTGTGAGCCACCACACCTGGCCTGTC
AGGACATCTTCACCTGTTCCTGTGTCCAAGATGGTATTGCCTAGGTTGTCTTCTGGGRTTTTTATAGCTTTGGGTTTTA
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CCAGCTATCTCAGCATCATTTATTGAATAGGGAATCTTTTCTCCAATGCTTGTTTTTGTTAGGTTTGTCACAGATCAGA
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GCATTCCTGATTTGGCTCTCAGCTTGACTATTGTTGGTGTGTAGGAATGCTAGTTATTTTTGCACATTAATTTCATATC
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CTAGCCAGGACTTCCAAAATGTTGAATAGGAAAGGCGAGAGAGGGCATCCTTGTCTTGTGCCCATTTTCAATGGTAATA
CTTCCAGCTTCTGCCCATTCAGTATGATGTTGACTGTGGGTATATCATTGATGGCTCTTATTTTTTGAGGGCTGTTCCT
TCAATACCTAGTTTATTGAGAGTTTTTAACATGAAGCGATCTTGAATTTTATCAAAGGCATTTTCCACATCTTTTGAGA
TAATCATGTGTTTTTTGTCTTTAGTTCTGTTTATGTGATGAATCACATTTATTGATTTGCATATGTTGAACCAACCTTG
CATCTCAAGGATGAAGCCTACTTGATTGTGGTGGATAAGCTTTTTGATGTGATACTGGATTTGGTTTGCCAGTATTTTA
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GAGAGTATGTATGTCCAGGAATTTATCCATTTCTTCTAGTTTTTCTAGTTTATGTGCATAAAGGTGTTCATAATACTCT
CTGATGGTTGTTTGTATTTCTGTGGGGTCAGTGGTAATATCCCCCTTATCATTTCTGATTGTGTTTATTGAATCTTCT
CTCTTTTCTTCTTTATTAATCTAGCTGGTGGTCTATGTAGTTTATTAATTTTTTCAAAAAAAGAACAACTCCTGGATTC
ACTGATCTTTTGAATGGTTTTCTGTGTCTCAATATTCTTCAGTTCAGCTCTGATTTTGGTTATTTCTTGTCATCTGCTG
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TTTTGATTTGGGCATTTCATGCTATAAATGTCTGTCCTAACACTGCCTTAGCTATTTTCCAGAGGTTCTGGTATGTTGT
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TCAGGAGAAAAGGCATTCAATTTCCATGTAATTGCATGGTTTTGAGTGAATTTCTTAGCCTTAGCTTCTAATTTGATTG
CACTCTGGTCTGAGAGATTGTTCATTATTATTTCAGTTCTTTTGCATTTGCTGAGTAGTGTTTTACTTCTGATTATGTG
ATCAATTTTAGAGTATGTGGCATGTGGCAATGAGAAGAATGTATATTCTGTTGTTTTKGGGTGGAGACTTCTGTAGATA
TCTATCAGATCCATTTGTTCCAGTGCTAAGTTCAAGTCCTGAATATCTTAATTTCCTGTCTTGATGATATATCTAATAT
TTTCAGTGATATGTTAAAGTCTCCTGCTATTATTGTGTGGGAGTCTAGGTCTCTTTGGAGGTCTCTAAGAACTTGCTTT
ATGAATCTGAGTGTTTCTCTCTGTTGGGTGTGTATATATTTAGGATAATTAGATCTTCTTGTTGAATTGAACCCTTTAC
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Fig. 6.265

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CATGATGTAATGCCCTTCTTTGTCTTTTTTATCTTTGTTGGTTTAAAGTCTTTTTTGTCAGAAACTAGGATTGCAACCC
CTGCTTTTTTCTGTTTTTCATTTGCTTGGTAGATTTTCCTCCATCCCTTTATTTTGAGCCTATGTGTATCATTGCATGT
GAGATGGGTCTCTTGAAGACAGCATACCAGTGAGTCTCGATTCTTTATCCAGCTTGCCACTCTGTGTCTTTTAATTGAG
GCATTTAGTCCATTTACATTTAAGGTTAATATTGTTATGTGTGAATTTGATTCTGTCATCATGATGTTAGCTGGTTATT
TTGCAGACTTGTTTATATGGTTGTTTTATAGAGTCATTGGTCTGTACACTTCAGTGTGTTTTTGTAGTGTCTGGTAATG
GTCTTTCCTTTCTATATTTAGTGTTTCTTTCAGGAGTTCTTGTAAGGCATGTCTGGTGGTAACAAATTCCCTCAGTATT
TGCTTGTCTGAACAGGATATTATTTCCCCTTCACTTATGATGCTTAGTTTGGCCAGATATGAAATTGTTGGTTGGAATT
TCTTTTTAAGAATTGTGAATATTGGCCCTCAATCTTTTCTGGCTTGTAGAGTTTCTGCTGAGAGGTCCACAGTTAGTCT
GATGGGCTTCCTTTTGTAGGTGACCTTACCTTTCTAGCTGCCTTTAACATGTTTTCTTTGATTTCAACCTTGGAGAATC
TAATGATTGTGTGTCTTGGGGATGACCTTCTTGTGAAGTACCTTACCAGGAGTTTCTGCATTTCCTGAATTTAAATGTT
GGCCTCTCTAACTAGGTTAGGGAAGTTCTCATGGATAACACTCTGAAATATGTTTTCCAAGTTGGCTTCATTCTCCTCA
TGTATTTCAGGGACACCAATGAGTCGTAGATTCAGTCTCTTTACATAATCTCATATTTCTCGGTTTTGTTCATTCCTTT
TCATTCTGTTTTCTCTATTCTTGTCTGACTGTCTTATTTTAGAAAGCCAGTTTTCAAGCTCTGAGATTTCTGAGATTTT
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TTGGGTTTCAACATACTCCTGTACTTCAATGATCTTCATTCCAATCCATATTTTGAATTCTATTTCTGTCATTTCAGCC
ATCTCAGCCTGGTTTAGAAGCTTGCTTTAGAAGGGACRCGGTTGTTTGGAGGAAAAAAGGCACTCTGGCCTTTTGAGTT
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GTTTTTTTTTTCCTTTTATCCTATTTGATGACCTTGAGGGTTTGATTGTGGTATTAGGTGGATTCAGCTGATTGGCTTC
ATTTCTGGAAGATTTTAGGGGTCCAACACTCAGCTCCCAACTTCTGGACTGTGTGCACTAACTCTGGGGGACTTGTATG
AGGCCCAAACTTTGTCTTCTCAYTCTTCAAGTTTTGGAATCCACTCAGCTAGGGGTGCTGAGATGGGACAGCTGCAGTG
AAGTGCTAGTGGGTGTGGGGGTGCCTGCCTCCTTGCAGATGTTCACCAGAGTGGCAGAGGCAATGCAGCTTACAGAGGT
GGGCAGGAGGCCCTTGCTGGAGCCTGTGTGCACAGTCACACTGGAGGTGGTGTTGGCTGGGGGTAGGGTGGTGGCAGGC
ACAGGTCTGAATGCCTTCTCTGTGCCCCACAAGCAGGAGTGATAGCTCAGGGTAGGGGAGGATTCACTGTTCTCTGTAC
AGCCCACGTGTTTACTTTTCAAATAATTTGATGAACTGAGTGAAAATATTGAATTTAGACTAGCTGATTTCATTAAACT
ATGCTTAAAACTTCAATTTATACAATTTTTTTCTTCATTTTAAAACTTTGAACCAAATTGGCTTGTAACATAAACTAAT
AGATTCAATTAGTTATTTGGAATTATTTTCAAAGATGTGTCCAACATTCTAGAGTTTAAAGTTTATATCCTCAAATTAC
AATTGCYTGTAAAGTTTACTGGTTTATTTCCTCAAATGATCATTGCTGGTTTATGTATATGAAATATTCTCAAACATAT
CAAGGTACAGATAAAGCTATGAGACACGTAAGCTTCTTCTGTTGTCCATTGGGTAGGTAGAGAAATGGATGACTTGGCT
TATTTAAGCTAATTACTAAAGGAAATAGAGAAGCCTTGAGTCCTTTCTGGAACAAAACAGAATTAAATATTCATTACAT
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TGAGCATATACCACTCTTTTTCTTTACAATACTCCTTTAGAGAAAAGAAGACTGCAGTTTGCATTTTGCCCTGGGTAAG
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GTGTCTCACCAGATAACCTATAGTATATTCATACCSAAACTTATCTTTTTCTAATACTTTAGTTATTAGTTTTTAGGTA
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TACTGGGAAAAAGGAGGATTCGAGTGGGGGAAAGTTCACGACATTGTCTCTCTGCCTCTAATCTTTCCTGCTCAAGTCC
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GAGCMGAGGCTGCTTACACTTTTTATAGGCCAATATCAATCATAACGTTTAAGTGTTTAATAAGTATCAATATTTATCT
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TTGTTGGCCAAACAGCAGTGAATCAACACATGTCTCTCCTCTGAAAGAATTTAATTTTTTGTGGGGAGGATAGTACAAAT
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GCAGTGGAGATGGCGAGAAGTGATCAGATTCTGGCAATGTTTTGAAACCAGAGCTAATTGTGTTTGATGACGGATTTTT
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AGTCCTGTTTGTGGAAAATGCTTGGAGTGCTGCCCCATTTCCCTCCACCCACTATAGAGTTCACCTGAGGCTATGGCAG
ACTCAGTCAGTTTCCCATCTCAGGCACCTGGGGGCTTTCTGCAGTACAAAGGAGCCTGCTGAGACCGTTTACAGAATTT
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Fig. 6.26G

TTTAACATTCCCAGTGGCAACACTCCAACATTGAGGGTCCAGAGTTGGTGCTCATATACCCCAGGTTCCACATTCTCCA
GTGGGACAGTCCTGAGGCATGTGTATATATGGTCCCACAAAAGGTCCCTGGGGAGATTGGGCCC``TCTGCCCCTGCTT
ATTAATACACACTGTATTAACTTTTCTTCCTTTGCTATCCCACTTCTCCATCACCTAACTTTGATTTCCTAGAATCGCC
TTCCTGCCTCTTCTCCAAATAAACTACGCACTCCAATATTTTGCAGTCTCCCTGGAAGGAATTAAACCTAAGACTATTA
AATACCCATATGTCAATCTCCAAGAAGGAGATTGATATACTATTCTGAAAACCAGGCAAAATGTCAGGGATGGAGATAC
AGTTGGGAGGCGTTAGTCTTTTTCACAAGCCTATTCAAAAGCCCATTGCACAAGCACTAGATATGTGTAGATTCCTACA
CAGACTTGTTACCGTCCTTGTCTGCCTCATGTCATCTCTTCCACATGAATACAATGATACTCATCAATGAATGCTCTGA
ACTTTCAGCAAAGGCCACAGTAGAATTTCTAGATAGTAGTAGATTGAGAAGTGATGACAATTTGATTGGAAGCTAAGGG
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GGTTTGGAAAACCTTGTAGCGGTTATGGAGGGGCACCCCAACCCATCCTTTGGTGTTGCCATTGCTTTTTAATTATGAT
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GAGTAGCAATGAATGAGCTCCTCAATAAATACATAGCACAGACACTGACAGGCAAAGTGAAGTTTGTAATCACTTCCCA
ATCAATTGTTCAACTGAATGTGAGAAGAAATCATTTTCTCTCCTGTTTTGCA_GAAGTATCCAAAATTTGGTTTTTCCTGCTGAG
CAGTTATCCAGATAATTGCAAATCAGTGGAATCATTTACATGAATAAAGATTTTTATTCTAATTAACGTGCTAAATCAA
GCACACCAAATGTCTCTAATTCTTATATTACTCAAATGGCAATATTTTTGTCAGTGACTATCATATGAAATTTCAGGTG
CCTAATGCATTTTCAAATGGCACTCATAAGCATACTAACGTTTTATAAATAAATGTCTCCATATTTTTAAGATGAATGG
ACAATTTTTGATTTTTAATRTAGTAGTCCCCAATTTTCAATTTTTCAATTTTCATTTACCGAATGGAGAAACAGCATTCATC
AGAGCAAAAGAGCAAGTTTCACAAGTCATGGGAACAAACATAGATGCTAATTTCCTATTTTAGGAAGTTCTTTTTAAAG
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Fig. 6.26-7

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Fig. 6.262

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ATGCGCAAGTAATTGAATGCCTATGTGGATTACGTTTAACTTATCCCACTCTCAAATTTGGGAGTTTCATTTTTATTCA
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Fig. 6.269

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GAGGAGAACTTTGGCTCTAGAGCCAAACAGACCCAGCTTCAGTGCCAACTCTGCTGCTTTCTTGATTGGGCAGGTTTCC
TAACCTCTCTGAGCCATTTCCATGTACATAAATAGAAAATTAAAATATCTACTTCATAAGGTCGTTGTGCAGATTAGAT
GAAAGCATGTACGTGAATGCAGGACTACCTGCCAACTCTTTCATAGCATTTTCCAAGTCTTCACTAACATTTCACAAAT
TTTGTGAAGCTTCTTTTGTGAAATACATTTTCAGATAAACTATCATCATTGGTACCACTCTTGGTCACAGAATAGTACA
ATAGAGCTGGCTTATAAATGGTTACCTTATAAGCAACTTGCACATACAACCCAGGCTCACCTTGGAAATGTAGTGTGCA
GGAATGATTGACAGCTGGCTAGCTCTACAGCCACACTGCTGCTGAGGTTCACATTTTCACATTATCACTTATAGCTATG
AGACTTTGGGCAAGTTACTTCATGTGCCTCAGTTTCCTCATTTGTAAAATGGAGTATTATGAGGATTAAGTGTGTTAGC

Fig. 6.27

CATGTAAAGTGCCTGGCACATAGACAACACTAAAATAAATGTCAGCTATTATTAATAATAGAATGTAAGCTCCACATGG
CAGCAATTTTTGTCCTTTTTTCCACCTTTGTATGGCCAGTGCCTGGAAGAGTGTGTGGTACATATTAGGTGCTTGGCGT
TTGTGTATTGAATGAATGAACCTATTAGCAAATGTGTTCCGATTCATGGCAAGTGTGGAGGATACTCTGCTTCCTTTTGC
CACATTCATGGGAAGAGGGGGGTTAGAGCTGAAGGAACCTGATGGTTCATTTGTTCAAAGTCATTTGTCTTACATTGAAG
AGCTTGAAGACCTTGAAAGGCCCCAGGGTCATGCAACTTGATAGCGGCCTGGTCAGAGCTAGAACCCTGGCTCTGTGTG
CTTCTGTGAAAATCTTTGTTCCAATGTCCCACCCCTCACTTCAATACATATGCCATCTAATTCARTGCTTTCAGTAAGG
CAGCAAAATGAATGAATGATTCAGAGCACACTTCTTGCAGACAGGAGAGTGTATAAGAGCCATAATTGGGCAATGACCA
TGGAAGGACAGGTTTCTGAATAAGAGAAAGACTTATCCAAGTAAAATTATGTTCAGACATTTTACTTACAAGTCTAAAA
ATCACTTGCTACAACAGCTACAATAGGGTTTTTGTAAGGAAGTACCACTTTGTAAACAACATTTTTGTAGAATGCATTA
TATGCATTTTTTTCAAAATTGAAGGAAAGCCCCATCTCAAACATTCAAAAGAATCYGCTCAATTATCAATGTAGACTAG
AAACAGAGATAGAGAGGGGAGATACAGTTTGAGTGTTTTCTGTATTATCATGACAGCACATGGAACAACCACAGGCAGA
AAGATTACAAACTCTGCAATAAATATAAAGGTATTGGTTTTATAAGGTGGCCTGCAGGCTACGGACCAGCACAGAGCTC
TGGCAAATGTGGAATGCCAGGAGATGTACAAAAAAATCAAGGTGACAAAAGAAGAAAAGCACTCCAAACTGCCAAGGG
TTTATAATCATGCTGTGGATATTTCTGATCATCAGTGTGTTAATGGTTTTCACTATAGTTAAAACTATACCATTTTTGA
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TTGACTGAGCAGCTGGATCAAAGCCTGTCTGATCTGGGCAGAGCAAATAAGGAATACTTCTTACAAAAGAGGAAGCATT
CAATTCTATACACTTTACCTTGGTACTCCTTGTTCTGGAGGCCTCAGCTTCTATCATGTTCTTTGCCACAGTCATATAC
CCACCTTCTACCTTGGCTCTCCCTCCTCCTAACTTCCATTATTTTTTCTACTTTCAAACTATAATTGACAAGCCTTTTT
AATTTCAACAAGGGTCTGAAGAGAGGTGTCTGATTTTCCGTAAGGCTGGGTTATATAAGCACACAGAGTTGAATCAGAG
CAAAGGAAAAATCCATTGGCAAAGGAAGAATACTATATGCTACTTTAAAACTTGTCTGAATGTATTATTTTATTTACTT
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TAGGCTCAAGGGATCCTCCAGCCACAGCCTCCTGGGTAGCTAGGACTACAGGTGCACCCTACCATTCCTGGCTTAAAAA
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CAATCAAATGCAAAGGACAGTTCAAAGGGGACAAGTATGTAAGCCACATTCCAAAAAAAAAAAAAAAAGCTTTGAAGTGA
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GAATTAAACATTCAAAAATGAACAGAACAGAATGTAAAAGATTATTCCTTCAGGTGGTAAATGTTCACAACCAGCTCTC
TATGGGGGAGGGGAAGGGCACTATTTGTAGTATTTGCTGATTTCTGTGCCAAAAAATTCTCTCACCTGCCCATTTCAAA
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CATCAAGTTTACATTCTGGTAGAGAAAGACAGAAAATAAACAAGTTAACAATAAATAAAATAAATAGAGACTGAGATGA
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TGATCCCTTTTTGGCAATTAACTTGCCTTTAATTGATTTGTGGATTACCCACTGGAAGTTGCTATGGATAAAGGAATGT
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TCAGGAATTTTTCTTTTCACATAATGGCAGTGTTTAGATAAAGAATGGAAAGTTAATGAATCAGATTTGTGTTCCCATC
CAAAGGGAAGATTTATGTCCTGTTTAAGAATCACATTAATTTGATGACGTTAGTACATTCTCTAGTGAAAGAGTGGCCA
CTTTAGTGGAGGAAAAAAACAAGCAACAAAACTCCTTCTGCCATCTTCAGGTTGTACCTGACGAAAAGCTTTTATTTGT
GGGTTCTATGAAGTATCACTGCCCTGATCTCAGATGAACTAAAAGATGAAAACATTTCATCGCTCAAATAATTGTTTAT
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CTGTGTGGAAGATGACAAGGACGGCTTTGTGCAACTTCTGTTGTATGCGCTGCTTTTCAACCCCTTGGATGAGATACTC
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GTGTGAATGTGCCAGTAAGAGAGAAAAAAAATGCTTAGCCATATTTACTCATATAACCAACATCTAGGTCAAAATCAAAA
TATTACCAGCTCCCAAGGGGACCCCTTTATTCTTCTTTCCAGTCACTATGCACCCGAAGATAACCACTAACTGACTTCC
AATAGCATAGATTTGACTACTTTTGAATTTTATATGAATGAAATTATACAATATAAACTCTTATTTTATTTCAGTCAAT
ATTGTGTTTATGGGATTCATTCATAGTGTTTTGTGCATTCTTTATTCTCATTTTGTTAGTATTGCACCAAATGAATATG

Fig. 6.27i

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CTACAATTTACTTATTCTAGTGTTGATGGATAGTTAAGTATTCCAGTTTTAGGCCATTATGAATTAGAGCCATTATGAA
CATTATTGGATATATCACTTAGTAAACAAATGTACATGTAGTGTTGCAGTGTTTTGTGCATACACGTAGGTTGGGTATA
TCCCAAGAGGTAGGATTTCTGGGTCATAGGTTATGCCTTTGTCTTTAGATATTGCCAAATGTGTTTCTATACTGGTTGC
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CTAGTAAATGCATAGGCAGTGTAAGTCACGCTAAAATAAATGTGTATTGCTTTTTCAAGATTTTGTTTACCCATGCATT
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GCACTATAAATGTATGCAATGTTTGTCAGTTAAATATTTTTAAATGAATTTTTTGAAAAGAGGAGGGGTGTGAAAATGA
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CAAAGGAGATAGGTAATTGTCTTCTCTCAAGAGCCACTAGACCACAAAATATCAACATCAAAGATGAATGCAAGCGCTA
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GGTGTGGACTATTATAAGGGACTCTGATGCTTCATGACAGGGAGTAATTTGATCCAAAGTACAACGGAGCTSTCGTGTG
GATTTAAGTTACCAAACTATTGAAGGGGACCCATGCACCACCAAGTATTCAAATCACAATATAATTTCATTATTCTCTAC
```

Fig. 6.272

CCTCCCTTGCAAATGTAATAACTCTGAACTGATAAACACATTGTACCTGTGTAGAAAAATGCCACCCAGAAGGAAGCTG
ACTGCTAATCTAGGCCTGTGCTGTCCAACAAGAGAGCTAGTAGCCATAGAGCACTTGAGCTGTGGCTAATCCAAATCAG
ATGTGCTGGTAATGTAACATGCACAAAGGATTTCAAATAATAGTGTGAAAAAAAGTAAAAATTTTCTCATTTGTAATTT
TCATAATGATTACATGTTTAAATACTAATACTGTGTATAGGGTTAAATAAAACATATTTAAATTAATTTCACCTGGTTG
TATCTAAGTGTCCATCAGCCAATTAACGGATAAAGAAAATATGGTACATACACACAATGGAGTACTATTCAGCCACAAA
AAGGAATGAGATCCTGTCATCTGCAACAATATGGATGGAACTGGAGGTCATTATATTAAGTGAAATAAGCCACACACAG
AAAGACAAACTTTGTATGTTCTCATTTGTGGGGAGGTAAAAATGAAGACAATTGAWCTCATGGAGATAGAGTGGAAAGAT
GGTTATCAGAACCTAGGAAGGGTAGTGAGTGGCAGGGGTGTGGGAGTGGAGATGGTTAATGGGTACAAAAATATATTTA
GATAGAATGATTAAGATCTAGTATTTGATAGCACAAAAGACTGATTGCAGTCACAATAATTTATTGTACATTTTTTAAT
TAAAGAGTATAATTGGATTGTTTGTAACACAAAGGATAAATTATTGAGGTGATGGATACCTCATTTACCCTGATGTGAT
TATTAGGCATTGTATGCCTTATATCAAAATATCTCATGTACCCCATAAATATACCTACTATGTACCCACAAAAGTAAAA
AGTAAATACATTAATTTCACCTGGTTGTAAAAATAACTTTTTGTAATATGGCTTTTTAAAAATTTAAAATTACATGTTG
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GAGAGAATTAAAACATTAGCAAATAGGGGCTGTTGATGAAAATGAGATTCCAGAGAGGGTGAGAAGACTGAACACATTC
CTGTAAAGCAACAGGGATTAATGCACCAATGAGAATTGCTTTTTTTTTTTTTAAACAAAACAAAACAAAACAACAAAAAA
ACCCACTAAATTTTTCTAGGGGAAAAGTAAATGCTAGAGTGATATTAGTGAATTAGGGGATTTGTGAAGATGCATTTGA
ATGTCAAGAATATAAATGTAGTTTTCTTAGTATTTTTGAATTCAGTGACCTTTTGTTAACCTCAAGAGACTGAGGCTAAG
ACACCAACATTTACCATGTGCTTCAATCATTCTCTCAAGGACGCACAGCTCCTCTGAGCTGTAATAGGAATTCAGGTCT
GTGTGACTCCTGAGCCCACATGACCGAACGTGTGCCCATGGAACACTGGTCCCTGCAACTGCTCTGCAAATAATGGTTC
TATACTTAAATCATTTTAAGAAATGTTGCATGTATCGTTACCGTCTTAAACATGATTCATTGATTTGCATATTAAAGGC
ACTGAGAATTGCTAAAAAACTGTTTAACTTTTTAAAATCTTTTTCTCAAACTTCTTTTGCCACAGAACACCTCCTCCCC
TGCACATGCATCTATATGCCATTTATCACCAGTTGCATGCTGAAACTAACCTTAGACAGAATGCTATTTGGAAATGCTG
AGATGCCTTAAATTGCCTTCTATCATCTCTCATTATTAGTTACTAAGGAAAATGGCTTTGAGAAAATATAAAATATTTC
AGGAAAATATGTAATTTGATGCTCAGATAACTCACTTTTTGCTACATCAGAAAAAGCAAAACTAGAATTTAAAAATAAA
CTTGGATGCATGTCTTGGTTATTCTTATATTCATAGTGTGTGATCCAAAGGGAGTCAAAGAAATCTGTAGCGAGGCTGC
TCACAGTCTAGGTCTCTTTTTGTGTAGGACAAAGGTAGGGCTTGGCTTCTAGTTGAAGCTACAGTTCTGTAGGACTTGG
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AGGTTTTTTCTCTAAATTATGCTAAAAATGTCATCGAGAAGCATAAAAACAGACTTTCATTAATAGTTGCTATAGCAGT
GGGAAATTTTTGTAGGAATTTAAGTAAAAAAGCAAGTTATTTAGCAGTTTTGTTCAAATTAGACATTTCTGCCATCAGC
CTGCCCCCTGGGAACTTGGAAAGTGCATACTTGCTCTTGAGTTCTGCTAGTGCCCTAACCACACGCCTTTCATGTCTAC
CATACTTCGATGATCTTGGATTCAAATATGATCCTCTGCCTTCCTCACAGAAAGAAACAATTTAGGGACATGGACTTAA
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CTTTTGCCTCTATAAAGCAGGGCTATGGTTATCTTTCCTTATTACCTCAGTTGATTGTTGGCTGTGAGAACCAAGTAAG
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TGAGTAAATAGATTCCTGACTCCCTAACAGGGCTATGTCTTGTGTCAGGAAGCCAGCGCACTGGGCTTCTTGGTAGGCA
GGAGACCCAAGTCATCATCTGCTACTAAGGCTGGGTCTTGGAAGATGGAGAAGGCCGTAGAAGAAAGGAAAGCCAAAAA
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TTGGTTATACCTCAAAACAATTCTCAACATTATTATCACTGCCACCACTTCTGTTAGCAGCTGTGAGAACAGAGGTCAT
GTGAGTAAGCAGTTTGCTAAGGGTTGCATAGCTTAAAAAGTAGTCAATAAGCTAAAGTTTGAACCCTGAGATGCCTGGC

Fig. 6.27.3

CTTCACAGCTAAACTTTTATTAATTACACCTGCTTCTAACAACAACAACCATGGCTAAAGTTTCATTAACCTACCAGTT
TCTAACAACATTAACAAAAACATTTTCAAAAATATAATAATAGTCATCAATAGGAATAAAGGGAAACAATCCAAAAATA
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AGCTTAAACTTCATGGAATAGTTTATTTGAAGCTTGTGGGTACCAATTCCAGGCAGCAAGATACTACTTTCAGAGATTC
AATTAACTTCCTTCAGAGTCACAGCAAAAAGCGATAGAGCTAAATTGATTACCTTGAAGTGCCTTAACCGTTTTGCTGC
TGTTGTTACTTCTTGAGCCAATTTATTCTCCCAGCAATGTCACTGTTACTGAAGTTGGTCTGTTCTTTATTCTCAGTTT
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TCCAAGCCATAGTTATTCCAAATGGCTGGGAGTGCCTTAGGACTTATTTCTGGAGTCTTTTGAACTATTTATATTATT
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CTTTTTAGTTTCTATCTTTAGGAATTTTCCGGTCCTTTCACAGAAAGACACTTTTTAAAGGAGCAATTCTTACTTCATC
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ACAGTGCCTCACACCTGTAATCCTAGCACTTTGGGAGGCTGAGGCGGGCAGATCACCTGAGGTCAGGAGTTTGAGACCA
GCCTGACCAAGGTGGTGAAACCCCATCTCTATAAAAATACAAAAATTAGCCAGGCATAATGGCAGGTTCCTGTAGTCCC
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CACTCCAGCCTGGGTGACAGAGCTAGGCTCCATCTCCAAAAAAAAAAAAAAAAAAAAAAAAAAAAAACTCCACTGGATCCT
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Fig. 6.274

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GAGTGACTGTGTTTTATTAGAATTGCAAGAAAGGGAACTTGAAAACATAAATTTAACATGGGGCAACAGGCCTTGGATT
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Fig. 6.275

AAAGAATAGTGCTAAAAAAAGGTGGAAGATGGTTAGGTGTAAAATTACCTCCTCTCACAGGACCTGGCACATGACAAAT
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CCAGTTACTTCAGGACTGTCTCATTTCATGATTAGTTGCCTTTGGTCTCATGCAGTGGTATCAGTTCATTCCTCTAGAG

Fig. 6.276

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TCTTACGCGAGGGAAAGAGCTGTTTTAAGTTGTTTATGAATAGAGAAAGCAGATTCAGTCAGACGTAATCTGACTCTGT

<center>Fig. 6.27</center>

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ATGCCTGCAGGGGTTGTGGCCTTTGTTAGTTTATTATGCCTGTAATTTGGGAAGTCATGCTGCAGAGATGTAAAGTGGG
ATGGGCCTGAAATAGGTCTTGTAATAAAAAGGCAGTTAGGGAAGCCTGCTGGTGTCCTGCCTGGTTACTACTATCTAAG
TGCAGCAATGTGTTAGTTGATTTTAATAAAGAGTTGAAATCATTGTTAACTGGCCTCCCTCCCACTGAGTGAAGAAGGC
AGATTACACAGGCACACGTTTRCCTCCCCTAGTGTGTACTATGAATGCAGCTGCTTCTGAAGGGCCAGCACGTGGCCTT
CTTAAAACTTTCCTTGGCTGGGGTTTTCCCTTCTTGCCAACTGGTGCCCACCAAGCCAGTGGATGCTCTCAACGTCTAG
GCAAATGCCAAAAATGTCATCTTTGGCTCAGATGTGCACTGGCCTGTTAGAGACAACTGCCATGCAGAGAAGGAATGTC
ATCAGCTGCTAGGTCTCAAATTATCAGCCTGGATTGACAGCTGCAGGATGTAGTGCCTGCCTATAGTACTGAGGCCATG
ACCCACCTCTGGAGAGACCACAGCCTCATCCCTAGAGTAGAACTGGAAATTCTCCATTGAGACAGGAAGGCAGCACTTG
AACCTGGCAGGTTTTGGAGTGAAGTGTCTAAGGACAGAATTTCTCATTTCTCCACCTCTGCCCATGTGTAAGGCAGCAA
TATTTTCATAGAGGTGAAGAAGGCAGGTGTCCCGTGGTTATTATGACATCTGTTTACCTCATTCTAGTCACTGTGCCAT
GTTTCACAGCCATTGTCAACTACATTTGGTGAAAACTGTTTCCTCCGTCCACAACTGGAACATTGACACTAACCACATT
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TCCATGTAAGCCAAGATAATATTTCCACTTCTCCTTCTGAACATGTGAGTACAGTTTATATATAAATCAACAGCAAGAT
GGAAATTTACTCTGAGCTTTGTGATGTTTTCACTTTAAGATGCTGTAGTTCTGATCCTTTATACATTTATTTCATTTTC
CCCATCTCAACCCCAGTATATAGCAACACTCACCTGCTCTAGAACATGATTATCAACTGTTTCAGCTATCAATGTAACA
TTAAAAAAAAAGTTCTATGTTCAAAGACTAAAGGAACCCAGGTAGTTTCTTTTAAACAGAAAGACTAGTTTTCATGATC
ATAAACATGTAAAGAAATATGTCATTTTTGAAATTTCATGAATCTTATGTCTATACCGATTCCAATTCCAAACTCAGAC
TGGCAATCTGCCTTTATTATTGCAAAGCCCCRTAGCTTGTTATCTTCATGTACCTCTTGATCAAGTATTTAAGTGAAAT
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GCAACTGGAAAGATTGAGTGTCCCGCCTGATTATTGGCTCAGGACTCTTATAAAGTTACTATTTCCTGATTCTTCCTGA
TCTCCTCCCATCATACTTCCTTTTGGGCCAGCTGTTGGCTAATCGCCGCGTGCTCAGTGACTTGCCAGTAATCTGGGAG
GGGCTGCATGCGCCATTTGGTGGTTGTAGTTATGCACATGTACTCTTTGGGCAATTTTCCTTTACTGGTCTAGTGCCCC
CAGAGGAAGGTCATATACCAGTCAAACTTTGCCATTTTGCCCCTTACTGTGCATGCCTGCTCAATTCCTAGGGTTTTAT
CGCAAAGTTGTTGCCTACAAGCTTAAGATGTTTCCTGTTAGGAAATTTCCCCCTCCCTGGTGCCAGCCATGACCATCTA
TCATTCCTGAAGAGGCCACCTGACAGTCACAATGACAGTCATTTGACTGTCTCCTGACATTCCTTGGGGCCCTATCCTG
CCCTGCTCATATCCACCTATCTACCTACCCTAACATATTCCCCCCTCAAGAGTGTGAGACTCAATTATTTGGGGCTAGT
GGATGAAGGTCAGTCCTCTGTGGCTGCTTCCTGCTGAAATAAGGGCTGTATTTGTCCTCTGGGTCTCAATCTCTTGCTA
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GCCATTATTGGATGGGGGTTCAACTGGGGGACCTTTTGGATCCCAGGGCAGTCCCTCTTCCAGTGGGACTTATCTCCAG
CTTATCACAAAGAGGGCAGGCTGTGTGGGCTTCCTCCCATTTGGCCCATTTGGAAAATTCCTTCTCCAGTAGCCTGGT
CTTCTGCATCAGTGGCAGTTACCTGGAGGAGGGTCCTTAGGAGACCCTGGAGGGGGCTGGTGGGCTTGTAAAGCAGCCA
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Fig. 6.278.

ATAGTTGAGCCTGCATCTTGTCCCTGCATTTCTCCTTTTCCTTAGCCCTGTCCTCCTTGTCCAGATCTCAGTTATAAAA
GACTGAGGAGGCTAATTTGAGGATTTGTGGCATAGGGGCACTGGGTTCTAAGGCTGACTTTTGTAAATTTTCCTAATGT
CTGGAGCAGCTTGGGTAAATTGTCCTTTAGGACTAGTCCCTCAGGAGATTCTGGGTTTAAGTTTGGTGTGCTTAATAAG
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GGCTTGATCTTACTTCATTTTATTACCTCTATTAGACACAAGAGCCTATGGTTCCTGGCCCATATTCCCGCTTTGGCAT
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Fig. 6.279

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GATTTCCAATTTCATCCATGTCCCTACAAAGGACATGAACTCATCATTTTTTATGGCTGCATAGCTTTCCATGGTGTAT
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Fig. 6.28c

CAGACAACCCACAGAGTGGGAGAAAAATCTTCACAATCTATACCTCTGACAAAGGGTAGTATCCAGAATCTACAAGGAC
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TGTTCCTGCCTTTCTGATTGTGGCTTTTAAGACCTTCCCCAGAGAGAGTACTGCCCTATACCCTGGGGGAAGGAATGCT
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GGAGGTTGGCACAGCCAGGTAAGGCATGGAAGCTCCACACCCCTTCCCCCATACCTCACCCTATATGCATCTCTTAATC
GGTAACATTTGCAATATCCTTTATAATAAACTAGCAAACATAAGTAAGTGTTCCCCTGGGTTCTGTGAGCCACTCCAGC
AAATTAATTGAACCCAAAGAGAGGATTATGAGTATGCCAACTTGGAGGTGGCCGGTTAGAAGCTCCAGAGGCCCACACT
TGTGACTGGTGTGGTGGGGGGCAGTCTTGGGAACTGAACCTTCAACCGGTGGGATCTGACATTATCTCCAGGTAGACAG
CGTTGGAACTGAATTACAGGACACCCAGCTAGTGTTGGCTGCTTGGTGTGGGGGGGAAACCTCACATGTTTGTTCGTAG
AAGGCTTCATCTGTGTTGATGATTTTTGTGGTGTGAGAGTAGAGGAAAAATGCCATCAGGGAGAGTTTTCTCTACACCC
TATAGCTCCTAGATGCTTTTTATTTAAGACATTCAAACCTTGTGAAAAACAAACAAACACAAAAATCTTTTTATTTAAG
ACGGTCAAACCTTGTTCAAAACAGCACAGACATTAACTCGGAATTTAGGATTTATTATTGATTAATTGACATTATCTTC
AGTCTGTTCATATTATTAAGTGATATAAGTCAGGTACACACACACATGTATGGGGAGTGTGTGGGGTGTCTGCATGAA
GTATACTGTCAGAAAATATACTCATTTCAAAGAAACACTGATTTAGGCCCTGGCAAATAAGGAAACATTTCTATTTCTT
CTAGAAATAACACATTCATTTGCCAACATCTGATCTACATATGACCTCTTAATACACACATGAAATAATAAAGTGTA
TTTTTCATTCATTTAATAAGCATGTGATTCACGTGACACTGTAAGCCTTTTTTTTTTTTTTTTTTTTTTTTGAGACA
GACCCTTTTTCTGTCGCCCAGGCTGGAGTGCAGTGGTGTGATCTCTGCTCACTGCAACCTCCACCTCCTGAGTTCAAGC
GATTCTCCTGCCTCAGCCTCCCAAGTAGCCGGGACTACAGGCACGGGACACCACGCCTGGCTAATTTTTTTGTATTTTT
TGTAGAGACAGGGTTTCGCCATGTTGGCCAGACTGGTCTGAAACTCCTGACCTCAGGTGATCTGCCCACCTCGGCCTCC
CAAAGTGCTGGGATTAAAGGCGTGAGCCACTGCGCCTGGGCGAGCATTTTTTTTTATTCCAGTGTTTCTCGTTAATCTA
CCCCATGGCGTTTTATTTCAGGTTGTATGTATGCCAGAGATGAAAAAAGGCAAATGACAGAAGAGCAAGTATATGGAGA
AAGATAGGAGGGACTTGAGAAAACTTGGCAATTTACTAAAATGAAATGGCATTGTTTCCATATTTACCAAAAACAGACA
AACAAACCTAATCATTCCTCTCATTTTGATCAGCTTTCATGTATATTTTCTAGGCCTAATCAAAATCTTCTTTGGTGTC
ACAAAATAATGGAAGAGACATGTATTGGTTTAGGCTCCTGCAGCAGTAGACCCCAATATAAGGAATATTAGTTGTTTAT
TTAGGAAATTCAAGAAACCTAAGTAAAGGAGTAGGGAAGTGAGACAGGAAATGGAAGGATCTCTGAAAAAATACACATC
ATCAAGTCCGTTCCCACAGAGGGAAACAGGAGCTCAGTCCCCATTGTTGGGTTCTGGAAGACAGTGTGGAACAGACCCC
AGAGTTATCCCAACTGTGAGATGAGGAAGCTGGGGTGCTTTTCACCAACTCTTCATCCATCACCTGCTTCCGGGGGTCA
TTAACAATCCACCACTCCCAGGTTATTTATGCTCAGGCCAGGCATGTAACTGCAGCCAGAAAATAGCCCTTGCCAAGAC
TCACAGGAATTAAAAACCTTCAGGCAAGGAGCTACTGGTCTTTGTAATAATAAGCTTTGAGAGGCGGGTATTCAGAGGC
TATGGGCACCAATAACTTCCCTTATGTGAGTCATAAACACAAAGTTAATTTTGTTTGCATGTGTATACAAGTATTTTGA
ATGAATGCTTCTCTTTGTTTTGTGTGTTTGGAAAATTTAAACAAAATTATTATTGTTTTTGTATTGGTATTTAAACATC
TGAATATGTAACATCAAAGATGATATTAAAAAGTATAAAGCAAATTATCACAAAATCAGATGTGCAGTTACATGTTTAC
TTGAAATTCTCTCTAGACTACATTTGAAATAGGAGCTTCCATAGGGGGAAATTTAAATTTTTTTTACAAGTTGTTTAAT
TATAATGTGCTTTAAAAACACTTTAGCATATCATAAACCTTCAAAACAAAATGAGTAATCATCTTCTGAGATCCATAGA
AAATGTTTCTTAAATGTCACTTTCAATTACCACTTCTGGAGCTAACTTGCAAAACAATTGTGTTTCCACCAACTGAAAT

Fig. 6.281

ATGTTTTCACCAAATTTAAGATAATCTGTGGTCAGTGTGAGCTTTGTCTTGATTTAGGAGGAGGAAAGCAAAGTTCCAT
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AGGTGAGAGGAAACTTTAAAAGTTACACTTATGGCCAGGCACGGTGGCTCACACCGTAATTCCAGCACTTTGGGAGACT
GAAATGGGTGGATCACTTGAGGCCAGGAGTTCAGGACCAGCCTGGTCAACATGGTGAAACCCCATCTCTACTAAAAATA
CAAAAATTAGCCTGGTGTGGTGGTGCATACCTGTAATCCCAGCCACTTGGCAGGCTGAGACATGAGAATCACTTGCACC
CAGGAGACAGAGGCTGCAGTGAGTGAGATCATCCCACTGCACTCCAACCTAGACAACAGAGTGACACTCTGTCTCAAAA
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CTTCTTTACATGGCCTCAAGTCTTTTCATCAAGATTTACTGATAGAATAACAGTAAATCCACATTACTCCAGGTGTCTC
TTTGCCCTTACCTAGCCTTTGTGTCTATGCCATTCTTCTCAAGGAAATTGCCTTCCTTTATCAGCTTTAAAAGTCCATA
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TTCATGATCTGTGTCATTCATTCATTCATTCATTTGCCCAATATTTCTTGAGTGCTTCCTAAGTTCAGGCACTGTTCTA
GGTACTGGGACATAGGAGGATAGAAGCCAGTCAAATTTTGCACTCACTGAAACTTTACTTTAGTAGGTAAGAATCTAT
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TTCACTACCCGTCCCTCTTTTTGTTGTAAACTCTCTAAGGCAAGTATAACCTCATCTATACATTAGTATTTATTTACCA
CAAGGACCCATAAGTATTTATTGATATATCTTCCTAAGTATGTGATAATATTCTAAATAGATAGTAGATTTAATCATTG
GGATTTATGTGAAGTTGGATTAAATAGATAACATGATAATTACATTTAGTATTGCATGGAAGGACACATTTTCTCACTT
ATAGAAGCCTAAACAAATGTCACAATGTTGATAGATTCCTTTATAGGATGTTAACTCTGAAGCTGTCGTGAAAATGTGT
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TGATCCCTCCTATTTCAATCTCATAAGACACTTATCTCTCCTCCAGAAAGATCTGAAGGGTTAAGTCACCTATTTATAC
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GCTATCATATGGTAGCTAACCTTGCCTAACCTTGCGAGTTTAACCTTGCTTTTTACGTGCATGCACGCGTGTGTGTGTG
TGTGTGTGTGTGTTACTAATTTCTAACTGTACCAGCTCCTGAATTAATATTTGGCAACATCAATATTCCTTGTATAA
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AATTATTCCTTATTTGTAACAACTACCTTACAGTGTTGTTTTGAAGATTAGAAGAGAATATATAGATGAAGTACCCAGT
GGACAGTAGGTATTGAATAAATGTTAGTTTCCTTCCACCTTCTCTGGCTTATGTAACAAAATTACTCATTCTACTAGTC
ACTCAGACTTTAGTTAATCATTAGCTAAAGAGTAACAATGCAGGATTTTTTTTTTAATAAATTGGCTAGGATTGGTTTT
ATGCCTGGATGGAATAGGCACTTGCCATTTCATCACAATTCAAAATTCCTCCTGAAGGCCTGAAGGCCTGAAATGTTTA
CCATTGGAAACTAATCCCACTGTATGGTTACTATTATTTATTATATCTATTAACTTATAAAATATTAAAATATAAAGTA
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CCAAACAATGGCTAAAAGGAAACCTAGTAACCACCCCGACCCCAAACTTGTTTGTTAGGCCAGTCAGATTTTTTTTTT
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TTATTATCCTGACACTGAGGACAAATATGAGTTGCTGTCCATCAGCATTCCAAAGCTGTTTTTTTTTTTGTTTTTTTTT
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AGAGCACTTTCTCTGAGAGCCGCCATTTTAACAGATCATTAAAGACACGATATTCACATGACGGTTGCTTACTCTCTGA
TGAAAACTACAAAAACAGAATACACAGGGAAGGTAATCTGAAGGTGATACCTTTTTCCTATGATCCTTGGCCTTATAAC
CACTAATCAAGCCTGAGGGCCGAAGTTCCTGCCTCATCTCTGCCTGATGTTACGAACTAGGCAGGCTAGCAGAGGCAAC
AGGCAACCTGGGCACCTGAAGAGCTACCTGGATAACCTAGAAGAACAGTAGGAGGTTAAAGATGAGGACAAGTCTATCA
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CATTTTCTCTAAGAGCATTCTATTAAACTGTTCCTAGTTATTAGCTTTCATAAAGGCACACAGAAAATGTTTCCTCTAC
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AACAAAGAGAGACAGTCTTAAGAGTTAACATCTATTGAGTCCGTGACTGCGTGCATTGTTCTTTCAAGTATTTTACCT
GCACTATTTAATGTAATTCTCCCAGCCACTCTGATTTCACTCATTTTACAATTGAGGAGACTGAGGCATAGAAAACTAA
AGCAATTTGGCTGGAGTAAGGGCCAGGTTGGGACTTATACCTTGGCGGTCTGCCTACAAGGTGTTCCTCACCACCATCT
GACACTGCTTTCTCTGTGTAGCCCAGAGTGTCAGCCTCAGTGCTTCAACTTGAGCTTTCAGGATCTATTTAAAGATGGA
AAATATAGTTACATTATGTCACCATTTGAGATGCAGAAAGACAGGGCCACCATTTTGCACAATTCTGAAAGCACAATTC
ATGCTGTGGTTTGTAAAAATGGTACTCCCTAGAGTTGGGCAATGGACAGCTCACACAGAGATGCAGTGGCCCTGCTAAA
CTAATGCCTTACATAAAAGAGTTTTACTCTTCATTCCCTTAAATGATTCTTAGTAACACTGGCTGACACACACACAAAA
CAGTTTCTAAGTAGGTCTGTCCGCATACTATGATCACTGGTCAACATTAAGTTTCCACAATTAATTTTACATAATCCAA
AGACTGCCAAGCAGTTTATCTGGGTAAACTAAACATTCTGTAGTCATTTATTCTGCTTCCAGTCATGCCCAGCAACAGG
TTGAGAAGACAAATGTTCTCAGAAATGATCTCCAAGGAGTTGGGAGCAGGCTGCTTATACGTCTAATTCACCAGAATAG
GTGAGCGTGGTCGTGGTGACCTTTTCATACTGTTGCAGAGCTGAGTATGAAGAGATGACTCACAGTCCTCCAATGCAAC

Fig. 6.282

ACAGGTGACCCTGCAGCCAGATTTTGCCTTCAGTGGTATGTGACTCCCATGGGGTCAGGAGAGTATCTCAGACATTGAA
ATTTGACCTAATGGCCTACCTACCTGCACACGCACCCTGCCCACTTACAGAAGGGCAAGAACTCTGCATTACTGAGGCC
TTGTTATTATTATAGTTCCTATTTAGGTAAGAACGCAATGGAGAAAAAATGCATTTGGTTATTGGGCCTCTGTTTGAAT
ATCCTGTGCTTTTTGCCAAACAATGCATTCTACATAATCTTAAAAAACAAAGTCCATTTCAAAGAACAAAAATAATGAC
CATATCCACTGAGCAATTGAGCAGAATGGGAATCGGAGTTTTAAACTCTGATATATCTGTTTTCCTTAGGGCTGAAATC
TTCTTTTCATGGTTCTAGTTTCTCTAATTGAAATAAGAACCTAACCCTGTTCAAAACTACATCTCTGGGAATGAGTGAA
AAATTAATGATCCTCCTATTTTGTTGGATCATAATAATGACTCTCATCCTGGTGGGGCCACCAGCAATGCTATTCTTCT
CAGACTCTATCTTAGTTCCTAAGCCACTCACCAGGTATTTAAAAGAATGATTTAACACAACTAGAATCATTTAAATAAC
TGTATTTTTTAAGAAAGCCATATTATTGCTTTAGTTTGGTCATTCATATATCCAAACCTTGGACTCATCATCTTGCAAT
CTCATGTGTTCAGTTATTCCAGAGAAAACATTAAGGAGAATTGTATTCTTCTTCCCAGCTAAATTTTAGGTCCTCAAAG
CTGGCAACCACAATTTATGCTTTCTAAAAATCATCTATGATACTAAGTATGAGCTGGATCATGAATAAAACCCTTGTAA
CAATAAATGCTTGATGACTTACTTCATCTCATTAGCAAGGGAAGGTAACTCATAATTATCAAGTACTACAGGAAATAG
GGCACTATCTGGAAATGCTAAATACATCTCCTTGCATGAATCACTTCTTACCAGAATTCCCTTTCTTTCCAGATCCCAC
CTCCAGGGTTCTGTGGTCCATACCAGGCACTACATTCCTCAGGGGCTACAGCCTATGAGCGTCTCATGGGCTATGAAA
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CAAAGCATAAGATTATGTCGACTTCAATAATTGTCAAATGAGTATTCTTAACATTTTACTAAATTAAAAAAACTTATGT
GCTGAGTTTTTTATTTTACAAGTATCTCCAAGTATGCTGGATGATTGCAAAGAAAATCAAGGCCAGTCATTGGTTAAAT
GAGTTTAATAGTAGCCACATAATTTCAAAAGCAAAATTATAAAGACCCTTCCCAGACTGTTGATAGCAAAAATAATCTA
CGTTGTGGAAAGTGGGTCCATGTTAATATGTTAGATATAAGTAGTGAGGCCTAAAAAGGTATTAAAACATCTTTGCTTA
AGGTACTACCTATTTGCAAGATTGTTATTTTAAAAATAGCTTATGTTTTAAATTGTTATTGCTTTTTATCACTCTAATA
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TTAAATCCTGCAGGAATCTGACTCACTGAAGACTGTCAACTGAATGAAGGCTTAAATTTCATGGTCTTGGGTGGGAGAA
CTTTTTCATATTCTTTCCCCATATGGAATAACAATCTGCCCTGAAAACAGGGAGTATTTTGGCATGATCTCTTTTTGCT
TATTTGCCTTCCATTTTCCATAAAGCAACTTTTGCCAAGCACCATACTTAAGACTCAACTTTTTTGCAAAAATATCAGA
CAAAGCACTGTCTTTAAGAACACAGAGAACACACTAGATCCCTTCTTCTGAAAATCACTGTTCTATGTTGTTTGTGGAT
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CAGGTTAAATAGTTAAGTCTGCTTTCCTATCCCTTATACAGAAGCTATCCATTCAAAAAAAAAAAAAAAAAAACTTCTTT
TTACTTTTTTTAATTTTTAATTTTGGTGCATACATAGTAGGTGTATATATTTGTGGAATACATACAAACATACAATGCA
TAATAATCACATCATGGTAAATGGGTATGCATCACCTCAAGCATTTATCCTTTTTTTATGTTACAAACAATCCAATTTT
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CAAACTGTTCTCCATAGTGGTTGTACTAATTTACATTCCCATCAAAAGTGTACCAGGGTTCCCTTTTCTTTACATCCTC
ACCAGCATTTCTTATTATTTGTCTTTTGCATAAAAGCCATTTTAGCTGGAGTGAGATGACATCTCACAATAGTTTTGAT
TTGCATTTCTCTGACGATCAATGATGTTGAGCACCTTTTCATATACTTGTTTACCATTTTTATGTCTTCTTTTGAGAAA
TGTGTGTTCAGATGTTTTGCCTATTTTTAAATCAGATTAATTTTTTCCTGTAGAGTTGTTTGAGCTTCTTATATATTCT
GATTATTAATCCCTTGTCAGATGGATAGTTTGCAAATATTTCCTCCCATTATGTGGGTTGTCTCTTCACTTTGTCAATT
GTTCCTTTGCTGTGCAGAAGCTTTTTAATTTGATGTGATCCTATTTGTCCCATTTGTCCATTTTTGCTTTGGTTGCCTA
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TTGGCGCCTTTGTTGAAAATGAGTTCGCTGTAGATGTATGGATTCATTTCTGGGTTCAAAACTGGTATTCCAGTTTTGT
TCTTTTTGCTCGGGATACCTTTGGCTATTTTGAGTCTTTTATAGTTTCATATAAATTTTAGGATTTTTTTTCTATTTCT
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TATTGATCCTTCCAATTCATGAATGTGGAATATCTTTCCATTTTTTTGTGTTCTCTTTTTTTCCATTAGTATTTGATAG
TTGATAGATCATTGTAGACTTTTTTGGTTTAGTTAACCCTAGGTACTTAATTTTATTTGTACCTATTGCAAATGGGATT
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CTACAACTTTGCTGAATTTATTTATCAGTTCTAATAGTGTTTTGGTGGAATCTTCAGGTTTTTCCAAATTTAAGATCAT
ATTTTCTGCAAATGGGATAATTTGCCTTCTTCCTTTCAAATTTCTTTCTTTGGTCTGATTGTTCTAGCAAGGACTTCCA
GTATTATTAATATATTGAATACTAGTGGTGAAAGTGGGCATCCTTATCTTGTCCCACCTCTTGGAAGAAGAGTCTTCAG
TTATACCCTATTCAGTATGATACCAGCTGTGAGTCTGTTATATATGACTTTATTGTGTTGAGGTATTTCCTTTTATAC
CCGATTTTTTGAGGATTTTTATCATGAAGGGATGTTGAATTTTATCAAATGCTTATTCAGCATCTGTTGAAATGGTCAT
ATGGTTTTTGTCCTTCCTGTCTTCCTTTTAGTAAAAGTGATTTTCTCTGGTGGCGTGTTTTAATTTTTTGCTTTTTATT
TTTTGAGAACCTGTTGTATATTTTTTGATTTGCGATTACCAGGAGGCTTGTAAATAATATATTGTAACTCATTATTTTA
AACTGATAACAACTTAACACTGATTGCATAAACAAACTAACAAGCAAAGGGAAAACCAATAAAAACTCTACATTTTAAC
TTTGTCCCCCGCTTTAAAAACTTTCTGTTGTCTTTATATCTTATTGTATTGTCTGTATTTCAAAAAATAGTTGTAGTTAT

Fig. 6.283

TATTTTTGATCAGTTTATCTTTTATTCTTTCTATTTA..GATATGAGTTGTTTACATACTATAATAACAGTGTTATAATA
TTCTGTGTTTTTCTGTGTATTTACTAATGCTAGTGAGCTTTGTTCCTCCAGATGATTTCTTATTGCTCATTAATATGCT
TTTCTTTCAGATTGAAGAGAATTCTTTAGCATTTCTTACAGGATAGGTCTGGTGTTGATGACATCCCTCAGCTTTTGTT
TGTTTGGGAAAGTCTTGATTTCTCCATGTTTGGAGAATATTTTTACAGGATATACTAGCCTGAGATAAAAGTTTTTTTC
CCTTTAGCACTGTGAATATGTCTCCTGCCCTATAAGGTTTCCACTGAGAAGTCTGCTGCCAGATATATTGGAACTCCAG
TGTATGTTTTTTCTTTATTTTTTCTTGCCGCTTTTCAGATTCTTTCTTATCTTTGACCTTTGTGAGTTTGAGTATTGAA
TGCCTTGAGCTAGTCTTCTTTGGGTTATATCTGCTTGGTGTTCTATAACCTTATTATACTTGAATATTGTTATATTTCT
CTAGGTTTAGGAAGGTCTCTGTTATTTTCCCTTTGAATAAATTTTCTACCCCCATCTCTCTTTCTAACTTTTTTTTAAG
GTCAATAACTCTTAGATTTGCCCTTTTGAGACTATTTTCTAAGTCTTCTAGGAATGCCTCTTTCTTTTTTATTATTTTT
TCTTTTGTCTTCTCTTTGTATTTTCAAGTAGGCTGTCTTCAAGCTCACAGTTCTTTCTTCTGCTTGATTGTTTTGTGCT
GTTAAGAGACTCTGAGACATTTTTCAGTATGTCAGTTGCGTGTTTTAACTCCAGAATTTCCGCTTGATTCTTTTAAATT
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CAGTCTGGGCTTGTTTGTACCTGTCCTTCTTGGTGAGGCTTTCCAGATGCTTGAAGAAACTTGGGTGTTGTGGTCTGAG
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CCTGGATCTGGGGGAGGGGTAACACAAATACCCCTGTGGCCACCACCACTGGGACTGCATAGGGTCAGACCTGAAGCCA
GCATAGCACTGGGTCTTGCCCAAAGCCTGCAGTAACCACTGCTTGGCTCCTGCCTATGTTTGCTCAAGGCACTAGGGCT
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GTCTAGAGATGACGTCCAGGAGCCAGGGCCTGGAGTCAGAAATCTTAGGAATCTACCGGTACTCTATTCTACGGTGGGT
GAGCTGGCACCCAAGTCACAAGACAAAGTCCTTCCCACTCTTCCTTCTCCTTTCCACAAGCAGAGGAGTCTCTCCCTGT
GGCCACCACTGCCCCAGGCCTGCGACAAGTACTGTCTGGCTACTGCTGATGTTCATTCAAGGCCCAGGGGCTCTTCAAT
CAGCTTGTGGTGAATGCTGCCAGGCCTAGGAATCTATTTTGGGGCAGTGGGCCCACCTGTAAGCCAGGGAAGGTCCAGA
AATACCATCTAAAGCCAAGCCCTGGAATCAGGAACCCCTAGAACCCCTTTGGTGCTTTACCCTGCTGTGGCTAAGCTGG
TACCTAAGCTGATTTTTGGTTCTTATGAAGGTGCATGAAGGTGCTTTTTTGTGTGGAGAGTTGTTCAATTTGTTGTTCC
TTTGGGAGGCTGCTATTTAGCCATATGGCTCCACCTCCCTCTCCATCCAAAAAGAAAACTTTAAATGAAAATATACATC
CAAAGAAAGTGTTTTTGAAATAAAAAATAAATGGTCTTAATTCCAAGGAAAGAACCAACATAGTAAAGTAAAAATGCAT
TTTTTCCAAAAGTTACTTGTTTGATCACATATTGTATTTTATTTTAATGAAATATCTAGAATAGGTAAGTCCATAGAGA
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AGTTACTTTTGAGTACACTTGCAGTCTGGTATCTAAATAATTGAAATATTTGAAAGTCGTGATTCCTAGAATAAAAACA
GTGACTAACTTTTAATGATTTCAATGTCATGTGCACAAAAAAACCATGTCCAGAGTAATAAGTTTACCCATTCTACCAT
TCTTTTCAAAGAATTGCTTAAATAAGTTTTAAAAAATTTCCTATTGAGATGTTACTTAAAATACGATTTGTTTAGTTCT
CAAGGTTATGCTTTCCCCAGGTTATGCTTCATTTGGGTAATCTAAGTTTAAGTCCTAATGTAAGACCTAGGGCATGGTA
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CACTTAGTGCCCTTCGGTCCTCTCATCCAGTACATGGATATACAGTCTTCATGTGGTTGATACAAATACGAATGTGCTA
ATGCATCTTTACCAAAAAGAGAGGTAAAAGAAAGGAAGGATAATCCTGCCAGAGATACATATGCAAAGATCTGAAAGTA
ATCGAGAGTGAGGGGTTGCAGATTGGACAACACTTAGTAGTGGCTCCAGATAAGGTTCTTAAAGACTGGCCAGGATGCA
GCCTTTTCATCCAGAGAAACAGCCATCTCATTCATAGACAGTGGTGACTGATTGGCAGATTTTATAAGATGACACAAAC
ACCCTCCCCATTGTCCTGCCAAAATGTGGGGAAACTCTTCCTGGAAAAGTTATTTCTTTGAATTGAGCTTCCAAAAGGT
TCCTATTAGAATTCAAATACTCTGTCTGGGTTGAATATCACATTGCATAATAGTTTCTCCAGATTAAGGCCATTTTTCC
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CCTTTTTGGCACAGAAGGCATCTAAAGTAACTTTTAGAGATAGAGGCTTATGAAAAACAAGAAAGAGGCAAGACTCAGT
TTTTGAAATCTAATTCCAGCCATGAAGCAAATGCCACAAAAGGGCACAGGAAGAAAATCTGTAAAGGGCTTATCTACCA
CCGTTGACCAAAGATTTATTCTGCTGTTAAGCAAATACCTTGTAAGCCAGATATTGTGCTAGGTATTCTGAATACAAAG
ATGCAGAAGCCTCATTCCCTGTTTTCTTTTTTTTTTTTTTTATTATACTTTAAGTTTTAGGGTACATGTGCACATTGTG
CAGGTTAGTTACATATGTATACATGTGCCATGCTGGTGCACTGCACCCACTAACGCGTCATCTAGCATTAGGTATATCT
CCCAATGCTATCCCTCCCCCCTCCCCCACCCCACCACAGTCCCCAGAGTGTGATATTCCCCTTCCTGTGTCCATGTGA
TCTCATTGTTCAATTCCCACCTATGAGTGAGAATATGCGGTGTTTGGTTTTTTGTTGCGATAGTTTACTGAGAATG
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ATATGTGCCACATTTTCTTAATCCAGTCTATCATTGTTGGACATTTGGGTTGGTTCCAAGTCTTTGCTATTGTGAATAA
TGCCGCAATAAACATACATGTGCATGTGTCTTTATAGCAGCATGATTTATAGTCATTTGGGTATATACCCAGTAATGGG
ATGGCTGGGTCAAATGGTATTTCTAGTTCTAGATCCCTGAGGAATCGCCACACTGACTTCCACAATGGTTGAACTAGTT
TACAGTCCCACCAACAGTGTAAAAGTGTTCCTATTTCTCCACATCCTCTCCAGCACCTGTTGTTTCCTGACTTTTTAAT
GATTGCCATTCTAACTGGTGTGAGATGATATCTCATAGTGGTTTTGATTTGCATTTCTCTGATGAACAGACACTTCTCA
AAAGAAGACATTTATGCAGCCAAAAAACACATGAAAAAATGCTCATCATCATTCCCTGTTTTTCAAAGAAACTCACAGCA

Fig. 6.28⁴

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TTATGGAGGCCCAGGTTTCTATTTGGGCCTCCATAATGCCTTGTGCTCTACCAAGAATGGTAAAACAATCCTTTCCAGT
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ACACATACATACATGCACACATACACATCATGAAGAGCAAGCTTTGGAATGAAATATACCTGGGTCCAAAAATTTTGAA
CAAATTGTTGACTTTCTTTATGTATGAAATTATTATTATCACATCTACTTTGGAGGGATGTTGTGATGTATAATTAAGA
```

Fig. 6.285

AAGTATAGTCTACTTAGCATGTAGGTAATTAATAATTAGTAGTTATTATTATTATCCTAACCTGACAGTTTATGAAAGA
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Fig. 6.286

ATCATACACATGAAACAAACTTGAATTTAACTATGTGGCTAAATGTATATCAAGTTGCTTATCATTCATTCATCTCTTA
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TGTCAAAGTTGGTTCAGGGTTTTTGAGCATGAGGTAAACTTTTTTTGGTCCTATTACCCAAGTTAGTGATGTATCTGTTA

Fig. 6.287

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TGAAATATACTTAAAGATATACAGCTTTTAGAAATGATTAAGATACCTTGAAATATTTGATGTATTTAAAACATAACTA
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GAAATTGAGCTCTTTTTAGTGAACTTGCTTGAGACATTCATATGGAATACAGAAAATAATAATTCCTAACATTCAGGAT
GTCATTCCATTGCAATGTGGCACAGAGTTTTTTCTTTGCATCAAGGAATTTATATGACTGATACTATTTTAGTCAGGAT
AAATTACTGTAAAATGTGTCCACTTTTAACCAATGGCATTCTAACATGGGATAGGCAACAGCGGGCTGTGATTTTTGCT
CTGATACTGCAACTTTGTTATAGAATAGCAAGTAACTGATTGGCAGCTAGATCACCAGCTAGCTGGAGAAAGCAGCTTC
TAGGTCTATAGTCATAGTTGGAGTCATAACCTCATTAATGATAGCTACAATGTATTGGGCATCTACTCTTTGACAGACA
ATGAACTTAGTACTATATACAGTTGAAGCAACTGTATGAATAAACTGTATGAATAAAACTTGAAGTCGCACAGCTTATA
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TGCTTTTCCTCACTATCTAAACTGCTTTGGCTACACATAGTTCTGTTTCATGGTATTAGAAAAGTAGTCAACAAGCTGC
TACTTATACTTTTCCTTCCTGTCCTAGACTGTTCAAGTTTCTTCTTTTAATCTTATGTATATCAAGAACATGTTTGCCT
ATGGTATTTGTCTGCCTTTCCCCCCAAGATTTGTATACAAGCCTCCAACCGTGAGCCCTAGAGATATATTGAGAAATAG
AACATGGACTTTGACACGGCTCCATTCCTTGGAGATCTAGCTAGTTCCTGACTCGTTCAGCTTATTGACTCTCTTCCTC
TGGTCAGACTATAAATTGCAGTTTTCCTTCTCTCTGGCATTTGACCAGTTAGTAGCACCATAAAAACCTTTGAAATAAAA
GAGACCACAGAGAATAACCTATAATCTGATTTTCCCCACCCCTAACCCCAATGTGGAGCTGGAAACATTTTGCTAGTCT
TTAACATTTCCAGTCAAGTAGGGTCAAGAAAAACTGTTAAAAACAATGTCAATGCTTAAGTGGTAAAAAGATTTGTTGA
AGAATATATATATATATATATTTAGAGACAGGGTCTCACTGTGTTTCCCAGGCTGAGGTACAGTGGCTATTTACAGG
TGTGATCATATGCACTACAGCCTTGAACTCCTAGTCTCAATCCTCCTGCCTTAGCCTTCCCAGTAACTGGGACTACAGG
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ATATAGCTAAGTAAGTAGCTGTAATTCGCATCTTGAAATTTTAAATTTACTTCTTTCCTCTCAATATAGTCTTCATCTG
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TCTCATTAGATCCTCATGATGAATCCTAGACAGTACTCATTAGTCTTATTCCTCAGACTGAGGCCTAATGAAGTAAAGT
GGCTTGCCTAAATTTGAGTGGCAAATAAGATAAATTTCATGATTCGCATATTTTCTTAATTATAAAATGTTACCTTTAGC
AATGATGTATATGTTAATATGTTCAATGAGTTGATCTACCTGAAGTGCTTAGCACATGGCTGGCTCCTTGTGAGTGACC
AGGAACTGTTAGCTTATATTATGCATTTTTGAAGTTTACACATTGGTATGACTTTCTAAATAATCAAGAACTCTAAATG
TGAAAGCTGATTCATTTTGTAAGGGTTTCTTTTTCCATAAGGTAGTCTTTTTTAACACAGTCAGAATTAAGCATTTATAT
```

Fig. 6.288.

AGAGCTAGGTATTCCAGATAAATCACGTTAATAGTGTGATTCAAAGAAGACACACTGCTATCTAGGTGGGTCAAAGAAG
TAAACTACTTAGTGGAGGTGGCAAAATAAGAAGTGGTCAGTGGGAAGCTAAACCACCTCTGATTTAGTTATTTAAAACT
CAATGAGAACCCTCTAAGGAAGCACAACTAGGCAGGGGTTTCATCTGAAAGGAATGGTCAGCTACCTGAGCATTATCTC
TAGGTGCACCTTTGTAAAGACTTTCTCCATGTATCTTTCCTACTGGGTGCCACCCTGTGGCTTGGTCCTACGGTAATCA
CACTGCCAGGCATTCTCTACCTCTACAAATCCCTTACTCTTCTAGTCCTAAGGTGCATTCCTGGTTGTGTAATTCACAG
ATTATTATCTCCCTGGAGCTGAGCCTCATCATCTCTATCAGACCAGACTGAGAAGGACTCCTTTTTCCTTTGTCCCGCC
ATCCCAGCTCAGTGCAGAAGGCTTAAGATGCTTTCTCCATCATTTCTAAAGAACAACAGACAAAAATAGAGATCTGGTT
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AGAGCAGGCATTTAGCATTCCTGTCTCTGATATCTACTATTTTATTATGAGTAGTTACCAAAACAAAACAAAACAAACT
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TGCTTGCCATTTAAATGGCCTTGATGATCTCATCATAACTCTATATATCTAATCTCACCATTAAATTTGCATGCAAATA
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CCTATGAAGAACATATAAATATTGTAAACAGAAAAACTCTTTTTGGATGTTGCAAATGCAAAGAAGAAAAATAAGTAAT
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TAAACTGTTGTCATTTTACAATAATTATTTTAGTATGATTTAATATCCAAATTCTGTTGCTTAGATTCACCCTCAGTGG
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ACATTTCTCTAACTCTTGAAGGATTACAATTATTGTCTTCTGTCTCACGTACCTCTCCTGTGAGCATGTTTTGGAGTAG
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CTTTTAGTAGATGGCACTTTTCAGGAGATTGTACTGGGTGAGTTGTAGACACAGTAGTTACTCTTTATGGCAAGAGTTG
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CTGCTCGAATGACTCTTCTCGTGTCATAAACTAGTAAAGAGTAGTGCATATAGAAATGCTGATAAATGTGTTGCACATG
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CTTCAAAGTCAGGGAACAACTTAGGGTTGGCTTACCTTCTTGTATTGTGAACTTTTCCGATTGTGAAGCCAGCTTTTAG
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TCCTTAATAATGAGGAAGTAATTTGTTTTAGTATTGTACAGAAACATTTTCTAGCTACCTGATCATTTAGTCTAGTTCT
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TATTGCTTTATTTAAGTGTTTATTAAGCTTGTTCTGCCACAGGTACCCTGCTGGCTGTTAGAGAGACAATGGTGAAATA
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GGGCACCATGCAAGAATGGAAAACCTGTTTACAACAAAATATCAAATTATTATCTGACTTTCACAATGAACCACATTAT
TCTATGACTGCCAGTCATACATCTGGACTACTATACCATTTGTGTGCAGTTACAGCAGCTCCAGTTCTATATGGCTAGT
TTTGAACTTAGTCTGACTTTAAAACCCATCCCATTTACTATCCTCTGCTTTTTTTTTTTTTTTGGTGTATATTGTGGGA
AATTCCATAAATGATTTTTATAAACTTTAGCCAGTAATCGGGAAAAACTGTAAGATTTTGTTTGTAAGCTTTATAAACG
TTACTTATAAGATTTTAATTGAATTCTAAATTAATTTCTAATTTGAACCTCTCCCTTCTCTTAGCTTTATATGCTTGGT
CATCCCTCCTTAAGTAATTAGCTTTTTTGAAGCTTTGTTTTTTTTGGTTTTTCACTGTTTCTGTTATATTGACGATTTG

Fig. 6.28g

ACTTTCTTATAAATTGTTAGTAATTTCCTGAATAAAGTACAGAAGTTTATCTTTCAAATTTCCCATGGATCATTTCTTA
ATAGTCAAACAATTGTCTAAATTTCCAATACACAAAGAAAAGTTTTAGCTAACTTACTTCAATAAGATGGATAATTTGT
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AGGAAATGTAACTACACATGCAATCTTGGGCTTTCCAAAATGTTTAAAGTACCAAATTCCCTTGGTGCCAATTGCTTTA
GGCATTTCATCTTCACTCAGTTGCATCCTAAATATAGCGTAGGTGTGACATACCCTGTAAATTTCAAATTCACATAAAG
AAGTCCCAGCCTTCGAGGCAAGTCTTAATTGAATACTAAATAAAAACCTTAGTGTCTTCTGAAATGTTTCTGAATAAAT
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ACCGACGGAGAGAAGAAATTATTATTTGCAATATGTGAATTTCTGACTAATAGGAGTAATAGAAAATAATTGGTCCCTT
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TAAATAGAACATTAATGCTACATGAAAGATACTTACTCTGGCATCCTCAAGAGAAAGTAGGTTCTTTTTTTTTCTCAGTT
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GTCCTGATGCTTTCCAATTTTTTTTTGACAACAGTTTTGCCTTTTTCAAATTCAAATGGTATAATTGGGGCTTGGTAGTT
GATGTTTATCTTAATTGAAACAGATTCTCTTCATCCTTTTGCTCTGAGACTCCCACTTTGAGGCTGAAAGGTCATTTTA
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CTTCCAGAGCTTACATAGAATAAACCATTTGGTTTTTGTCTGACTTTCTTTTAAAAGGAAGTTATTGTGCTTCCTTTTT
AATGCTCTAACTACAGAGAATGGGGAAATCCTGTTTTTCATTTATTCTTATCAGAAAGAAAGTGTATTTTGCTCAGTAAT
TGCAATAAAATACTACTGGGGTTACAGCTACTATACTTGTCAACAAATTTAAGGGATATTCTTTAGCTGCATTACTTTA
AGGATTTCTTGTCTATCTTTTATCTAGCAACTTCATTTTTTAACATTTACTCTATGTACTTGTGTAAAGGTGCAAAAAA
TATATGTACAAAAATGTTCTGCTGGAGTTTTTGTAACAGAACACTGGAGAAAGCCTAAATGTCAATCGGTATATGGCTG

Fig. 6.290

ATTGAACAACTGATGCACAACTATAAGACAATACTGTGCATCTATGAGGAAGACCTCTCTAGTATGACACTATCTCCAG
ATGACATTTACTATTGTATTATTTTTATACTTTGAAAAGAAGATATGCACACTCCCATGTATCCATTTTCACTAGTTTT
CTTTGTATATACTCTTAGAGTTTATGTAAGTGCAATGAAAACATAAATTCTATTCTCTTCCTCCCTTTTTAAACAAAAA
GAAGTATAATCTGCACATTCATTGTTCTGCAGTTTGTATTTTGCACTTGACAGTGTGTCAATACATAGAGAAGCTTCTT
GTGTTGTTCTGGAAGACCTTTGTTGAGATATAACTCACATACCCTACTCTTCATCAATACAGGGTGTATAATTTCGTGA
TTTTTAGTATATTCACACACTTTTGCAACCGTCACCAAAATCAATGTAGAACATTTTACTACCACTATAAGAAACCCCT
TACCTTATAGTTATCACCCTCCATCTCCCCCATTACCTGCTCCATATCACCTGCCTCCCTTCCACCCCACAGCTCTAGG
CAACGGCTAATCTACGTTTTTTTCCTCTATAATTTGCCTACTCTGAATGTTTCATATAAATGGAATTATATAATATGAG
GTCTTTTGTGACTTGCTTCTTTCACTTAGCATAATGTTTTCAAGTCTCATTCATGTTGTAGCATATATCAGATTTCATT
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TGTTTCACTTTTTGGCTATTTTGAAATGTGCTGCAATGAACATTTATGCAAAGATTTTGTGTGTGCATATATTTTCCT
TTCTTCTGGGTATATTCCTAGTAGTGGAATTGCTGGGACAAATGGTAGCTCTAAGTTTAACCTTTTGGTTATTTTCCAG
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ATGTATTCATTAAGCATTTATTTATTCAACACTCCTATGACATCACTCCTCACTGAGTGATGATTCCTTCTAGGGTGCC
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ATGATGATGATCTAATTGCAATGCTAAGAACCATGGCTGGGCTACGACTGTGTGTGCACGTTGTTATGCTGGGGAGCAA
GGGTACCTAGGAAAGAAAGCAATAATGAAAGTATCACTTGGGTATTTGTTTATTTCTGGATCTTATTCTATCTGCTCCA
TATTATCACTTTGGCATTTAAAATCTAGATGATGGCTGGGTGGGTGCAGTAGCTCACACCTATAATCCTAACACTCTGG
GAGGCCGAGGTGGGAGGATCACTTGAGTCCAGGAATGTAAGACCAGCCTGGGCAACATAGGAAGACCCTGTCTCTACAA
AAAAATATTTTTAAAAAATTCTCCAGGCGTGGTTGCACACACCTGTAGTCCCACCTACTCGAGAGGCTGAGGTGGGAGGA
TTGCTTGAGACTGGGAGGTCAAGGCTGCAGTGAGCTGTGATTGTGCCACTACACTCCAGCCTGGGTGACAGCAAGACCC
CATCTCGGGGAAAAAAAAACCCAGATGATATGGTTAACCATATTCAGTCATTATTCAGTTATTAGAAAATAAGATTTA
CAAGGCATCCATGGAGGGGAAAACAATTACACACCTGGTTAGTTGGGCTGAGGCTTGCAGAGATAAAATCACCTGCACA
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TGTTAATTTGCTAAGGATAATAGCCTCCAGGTCTATCCATGTTAAAAGACATGATCTCATTCTTTTTTATGGCGTCATG
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TGCCCAGTAATAGGATTGCATGGTCAAACGGTAGTTCTGCTCTTAGCTCTTTGAGGAATTGCCACACTGCTTTCCACAA
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TTTGTCTTTTTAATAATAGCCATTATGACTGGTATGAGATGGTATCTTGTGGTTTTGATTTGCATTTCTCTAATAATCA
GTGTTATTAAGGGTTTTTTCATATGCTTGTTGGCCGCATGTATGTCTTCTTTTGAAAAGTGTCTGTTCATGTCTTTTGC
CCACCTTTTAATGGGGTTGTTTGTTTTCCTCTTCTAAATTTGTTTAAGTTCCTTATAGATGCAGGATATTTGACCTTTG
CCAGATATATAGTTTGCAAATATTTTTCTCCCATTCTGTAGGTTGTCTGTTTAACTCTGTTGATAGTTTCTTTGGCTGTG

Fig. 6.291

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CAGAAGCTTTTAAGTTTATTTAGATCCCACTTGTCAATTTTTGCTTTTATTGTGATTGCTTTTGGTGTGTCTTTGTCATGA
ATCTTTGCCTGCTGCTATGTTCAGGATGGGATTGCCTAGGTCATCTTCCAGGGTTTCTATAGTTTTGGGTTTTATATTT
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GCCAGTTATCCCAGCACCATTTATTGAATAGGGAGTGTTTTCTCCATTGCTTTTCTTTAGCTTTGTCGAAGATCAGATG
GTCCTAGGTGTGCAGTTTTATTTCTGAGCTCTCTATTCTGATCCATTGGTCTATCTGGCTGTTCTTGTACCAGTACCAT
GCTGTTATGGTTACTGTATCCTTGTATTATAGTTTGAAGTTGGGCAAGGTGATGCCTCCAGCTTTGTTATTTTTATTA
GAATTGCCTTGGCTATTCGGGCTCTTTTTGGTTCCATATGAATTTTAAAAAGTTTTTTCTAGTTCTGTATTTCACAGCT
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TCTACAAAATATTTATATTCAAGGCACAATGACCCTTACATATTCTTTTTTTCTTTTCTTTCTTTCTTTTTTTTTTTTT
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CTTAAGCGATACTCCCACCTCAACCTCATGAGTAGCTGCAACAACAGGCGTGCCCCACCACTCCTGGCTTGTTTTTTTT
AATTTTTTTGTTTTTTTTTTTGTAGAGACAGTGTCTCCCTATGTTGCCCAAGCTGGATTCAAACAATCTGCCTGCCTC
AGCCTCCCAAAGTGCTTGGATTACAGGTATGAACCACCATGCCCAGCCCTTAAAATAGTTTTTATTATAAAGGCAATTT
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GCCAAAGTGAGAGGACTGCTTGAAGCCTAGAAATAGAGACCAGCCTGGCAACAAAGCAAGGCCCTGTCTCTACAAAAAA
TAAAAAATAAAAATTATTTTTTTATGGTGTGTCCCTGTAGTACCAGCTCTCAAGAGGCTGAGGTGGGAGGATTGTGTTC
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TATACACACACACAATACGACACGTATAGAAAAAATAAAAATACAAATATAAATTTCTCATAACCTCAACACACT
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AGGCTGTTTCTAGCTTTTCCCAATTATTAAAATTATCAGCAAGGAACATCTTTGAACTGAAACATTGGCTTATCTCTGA
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CTAAAGGTTATACCAGGTTACACTCCAAGCCACACTGCTTGAGAGTTCTCATCTTACTGTGCTGCTGCAGGAAATGTTT
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Fig. 6.292

TTCTTTCTAGACATTTATGTGTTTTTATTAGGATAATGAATACAGAATTATCAACATCATGATCATAGTTTTTTTGGTT
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Fig. 6.293

GAGAAAAGCCCTCTTAAATGCCTACTTATTTCAAGTCAAATTTAGTTAAACTACATAACTTAGTCCCCTGAGTGTCAGT
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Fig. 6.295

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TGAAGGGAATTAGGACAAGATTTTAAAGTAACCTAAGTGCCGTGATAAAGAATCCACTTTTATAGGCAAAAGTAGTTAT
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GCAGAAGATCAGTGTAACTACTACCATCCTTACGTCTTCTCAATGAGCTTCATTATTCCTCCCAGAGTGGCTAGAGTAG
GTAGTTGATGAGCATTTGTGGAATGGATAGGTATCTCTTTTCTATGACCTACTCTATCACCCTCAGGATCTATTAATCC
TTTGAGTTTGATCTCTAAGGTGGCAATGAAATATTTCTTACACCAGAAATACACTGAAACTCAGAGAAAGAGGGCTCAA
ATTTGAGGAGAGTGCTATACTGGCAAGAGGATGCCCCAGGTAATATCCTTTCCTTCTTCTAGAGAGGAACATTTGGCTC
TAGAAAGGCATTTAGCCTATGAGTTCAGAACAGGAATGGTCTTAGGCATCTGATTCAACCTACGACATTGCCAGTCTGA
ACCAACTCATCCTGATCCCACTGCAAATGAGCTAGGCAACAGTGAAGAGTGGTTCTAGTCTGGGTTTGGATAAATTGTG
AAATATTTCTAGAATTTTTCTATAGACTCTGAACCATTTCCCCAGAAGTGTGCACATTCAAAATGTCATTTTGCTTCAG
AATAACATTTTGAATAAGGCAAGCCTAATTTTTGTTTTTTAACAGAATTTTAAAATAATTAATTCACAATATTTGTACA
TAGGATAATGGGATATTTTGTTACATAAATAGGATGTATAATGACCGAGACAAATTATTTAGGGTATCCAGCACCTTGG
GCATTTATCATTTCTATGTAGTGGAATAATTTCAATTTCTTTCTTCTGGCTATTTTGAAATATATGATACATTATTCTT
AACTATAATCTCCCTATTCTGCTATCACACATTGTAACTTATTTCTTCTTCTATCCAGCTGTATGTTTGCATACGTTAGCCA
ACCTCTCTTCATTCCCCTCCTCCTGGTAGCTATCATTCTATTACCTTCATGATATCAACTTTTTTTAGCACCCACTTAT
GAGTGAGAACATGCCATATTTGTGTTTCTATGCCTGGCTTATTTCACTTAACATTCAGGTCAATTCATGTTGCTGCAAA
TGGCAGGATTTTGCTCTTTTTATGACTGCATAGTATTCCATTGTATGTGTATGTGTGTGTATATATATATATATATATG
TATTTTCTCTATCGGTTTGTTTGTTGGTGGATACTTTGGTTGATTCCATATCTTAGCTATTGTAGATAGTACTGCAATA
AACATAGGGGTTCAGGTATCCCTTTGATATACAGATTTCCTTTCCTTTCCTTTAGATAAACACCCAGTAGGGGGATTGC
TGGATTGTATAATAGTTCTACTTTTAGTTTTCTGGGAAATCTCTATGCTGTTTTCCATAATGGCCATATTAATTTATAT
TCCCACCAACTAAGAATTCCCTTTTCTCCACTTCCTCACCAGCATCTATTTTTTTTTTTTTTGGTCTTTTAATAATAGC
CATTCTAAGGTAAGATACCTCACTGTGGTTTTGATTTATAGTTCCCTGATGATTAGTAATGTTGAGTACTTTTTACATA
CCTGTTGGTCATTTTACATCTTCTTTTGAGAAATGTCTTCATGTCCTTTGTCCACTTTTTGATGGGATTATTTGGGTTT
GGTTTGGTTTTGTTTTTGCTGTTAACTTGAGTTCTTGGCATATTCTGGATATTAGTCCGTTGTCAGATATATATTTTGC
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Fig. 6.295

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AAATATTTTCTCCCATTTAACATGTTGTCTTTTCACTCTGTTGATTATTTTCTTTGCTATGTAGAGATTTTTGGTTTTA
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CTGAAATCTCATTGGCTCAAGTGAAGCACGGGTGTAAATCCTGAGTGTTTGTTAATCTCGGTTCTGTGTGTCTTCTGTA
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AGAACCAGACTGGCTTTCACTTTACACAACAATCACAACTGTGTGTAATGATTATAGACTTATTCTCTAACATCTCTTA
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AATAAGCCCCCAAATTAAATGACTTAAAAAATAAACTCATTCTTCATCAAGTAGCGTTCTAGGATGGTGTTTAGGAGGA
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CTTTAGCATCCAGCTGGCAGAAAAAGAACATCAAGATAGGCATGTGGGAGGAGCTATGAGCCAGTCCTGAAAGTGGCAT
GCAATATTTCTGCTCACATCTAGTCACTTGGCCAAATCTAATTGCAAGGGAACGAGAACTTCTTTTATGCCCAGGAACA
GGCAGAGAATGGAATATGGCAGACAACAAGCAGAATTTCCTATACTCACAGTCCTGATTATTGACTTCTATGCTTTCCC
AAGGTCATTTTGTCTTCAGCCAGATTCCATTTCAAGGAAACATGAAAATGTTTCTTCACTCTATAAAATCATTGTTGAA
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CCCAATATAAGAAATCATGAGAGAAAGCTAATCTCAGTGTACTTATTATACAATGAGACACATAATGAGACATATCTTT
TTTTATTCTCCACTATATATTTTAAAAGAATTGAAGAGGCAAGTGATTGTTTATGGCCATCGTAAGATAATATTCTTAT
CATTGCTTTGCCCAGTAAGATTGAAAAAATTGTCTGTAATTCAGGCCACACAACCCTCTGAAGCCCTTCGCCATTCATT
CAAGCCCCATTAAGGCAGAAAACCCATCTGTTTGGATTGTTGAGGTTGGAAACTGAATAATATCACTTCTCCAAAATAG
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GATTTATCATTCAATCATTCTTATTTAGTCAATAAAATCTTTTGGAGATTAATTTTTAGTAAGCCTAAATACCCAAATA
GCCAGGAATGTGATTGAGGATCACATTTTTAAAACCCATCCCTCAAAAAGAAAATTGTAATATCTTGAGAGACAGGTAT
GGTTTGAAGATCACCCCTCTTCAAAGTGAGTTCAATATCTGACCTAATGGAATCACTCCCCATTCCCCAGGACTAGGTG
ACACTCACTGATCTGGGAAAAATAAACACGTGCAACTAACAAGAGAAATTTTGAGAATTATGAGTAAGCTTTGAAAATT
TGGGTACTGAAAATAGAGAGGGAGGGAGAGAGGGAGGTGATAGGGAGGGAGGGAAAGAATAAAAAGGAATGAAGAAAAA
ACTAATAACAATTTATCTCTTAAGAAAATAGAAGGTGCAGCTGGGTACGGTGGCTCACGCCTATAATCCCAGCACTTTG
GGAGGCCGAGGCAGGCGGATCGCCTGAGGTCGGGAGTTCGAGACCAGCCTGACCAACATGGAGAAACCCCAAATACAAA
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TAAAATAAGTAAAAAACAAAGTTCATCAGAAGCTGTTAAATGCGATGGAGAAAATTAAGAGTAAGGAAGGTTTTTTGT
TTGTTTGTTCAAAAAAAAAAAAGGTTAGGAGGTTAGGGGTGCCAGTTTTAAATAGGCCAGTATGGGAATATCTCATGAA
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GATTAGAGAGGCACCAGGGACCAGGTCACTGAGGGCCTTAAACCATTGGGAGAACGTAGTATTTTCTCTGAATAAAATA
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CCTGACAGTTAGTATTTTTTTTCTGATCCTCTTCTCTCCTCCCACCCTCCATCCTCAAGTTGGCCCCAGTGTCTATTGT
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GCTGCATAGTTTTCCATGGTATATATGTACCAGATTTTTCTTTATCCAATCTATCATTCATGGACATTTAGTTTGATTC
CATGTCTTTGTGATTGTGAATAGGGCTGTAATTAACACATGCATGCACGTATCTTTATGGTAGAATGATTTATATTCCT
TTTGGTATATACCCTGTAGTTTGATTGCTGAGTTGAATGGTAGTTCTGTTTTTAGTTCTTTCAGGAATCACCACACTGC
TTTCCACAATGGTGAACTATTAATAATTTACATTCTAACCAGTAGTATATAAGCGTTTCCTTTTCTCTGCAACCTTGC
CAGCATCTGTTTATTTTTTGACTTTTTAGTAATAGCCATTCTGACTGGTATGAAATGGTATCTCACTGTGGTTTTGATTT
GTATTTCTTTAATGATTAGTGATATTGAGCATTTTTTCATACACTTATTGGCCATATGTATGCCTTCTTTTGAAAAATG
TTCATGTCCTTTGCCCACTTTTTAATGGGGCGGTTTGGTTTTTGCTTGAATGTTCGTTTAATTTCGTTGTACATTCTGG
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Fig. 6.296

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GTATTAGACTTTTGTCAGGTAATACAATTTACAAACATTTTCTCCCATTCTGTAGGTTATCTGTTTACTCTCTGTTAAT
AGTTGTGTTTTTTTTTCCTGTGCAGAAGCTCTTTAGTTTAATTAAGTTCCATTTGTCAATTTTTGCTTTTGTTGCAATT
GCTTTTGGCATTTTTATCATGAAATCTTTGCCAGTTCCTATGTTCAAAGTGGTATTTATTTGGTTATCTTCCAGGGTTT
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AGGTCCAGCCTGGCCTGGAGAGCATGTGTGCTTTTCTGCTCTTCCCTCCATACTCTTAGTTAGTGGTCAGCACATGATAAT
AGGATGTAAGGGAAGAAATAAATCTCTACTTGTTTTATTCTCAATCACAGTGACACCTGCATGAAATTGCCAGAGAAAT
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CAAGAGTCCAAATACCATGGCCTCATGTTAGGGCCATTTGAGATCCTTTTTGAAAAAGTTGCTTTATTGTGCTAAAAAC
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ACAGCATATCTCTAGAATTTATTCATCTTGCATGATTAAATGATACACATTGAACAACAACTCTCCATTTCTCCCTCCA
CCCAGGCCCAGCAACCACCATGCTACTTCCTGCTTCTATGAGTTCAACTACTTTAAATCCCTCATATAAGAGGGATCAT
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AGGATATCCTTCTTTTAAGGCTAAATAATATTCCATTGTATGTATATACCACATTTTTTAATCCATTCGTTCATTGATG
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CTGATCTCAATTTTTTGGATAAATACCCAGAAGTTGAATTGGTAGATCATATGGTAGTTCTGTTTTTAATTTTTTGAGG
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TGAAAAGAACAAAATATCTGGAGATCTCACACTTCCTGATTTCAAAACATATTACTAAGTGACAGTAATCAAAACAAT
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CATAACACTGGCCTAATAAATACTTTTCAGAAAAATTGGCCTTTTATCATATCATTGTGTCAAAAGTGTGCTAAGCAAG
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CTCCCCTTGAGGGCAAAGCATCAAAATGAGTTCCAATGTGTTGGCTAACAGCATGAGCTACAATACACAGGGGAGAGTG
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GAAGGACTGTTTCTGCTCATTATATTCAGCTGGAGCCCCACCTAGGTTGTTGGGACCTCCAAGCCTTCCACACTTCCTG
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AGCCATGTGTTCAAATAAAGTTAAAGAGAATGACCTAGATGAAGATCTCCCTGGGAGAAAAAGAGTAATAAAAATGAAA
AACAATAGCTGCTAAATAATGGGAGTGTACTAGCTGCCAGGCAATGTGGTGAATGCTTTCCAAACATTTTCTCTTCTAA
TCCTTATAATGGCCCTAGAGCTAGGTATTGTAATTATCTTTATCAGCAGTGAAAGCCTAGAAAATACAGCAACTTGCCC
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Fig. 6.29₹

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TAGAGCACCCAAATAGGAAGTGGCTGAACCAAGATTTCAAATTTAACCTGTGTTTTCTAACTCTTAAATAAACATCTTA
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CAGTTGAGAGCGTGGATGAAAGGTAGATTGGAACTGTATTACCATCCAGTAATTCCACTGAAATGGAACAGTGTGGAGA
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TGGGTCTGTGGAGTAAATGGTCATGAGTAGTGCTAGCCACACACATTCTTTAACACTTCTGAATGCAGCAGGCCAGCTG
GTCATTACTGTTACTGCCCCCTCATGGGAGACCAAATATAGCTATATCATTCCTCCTCCAAAACTGCCTTCTGGCAAAA
GTTATCTCTTTTAAGAATTGTGTGCATTCTTAAATTGCATTAAGGGAAGCAAACATATGGAACAAGAAAGCACCTCACT
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ATAACACCACGATGTGGACTATTGATAAAATTATGTTTCAAAGATAACTAGCTTATTTGAAGCCATCTATAATTTCAGC
CCAAACTGACTAATTAGGCTAACTTTCAGAAAGTTAAGTAAGTGTCTAATTCTCAGTTAACTTACTACCTACTGAGTAA
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TATTTAACCCTTCAGTAGAAAATAGTTTCCTTGTTTAAGAAAATGATCTCCATCTAGTGGACAAATTTTTATATTACAG
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CTGTTCATGCACAGGAGGGAGGGGGTATTTTCTTTTTTAAAGTGTAGCAATAAGAATATCTTGATATTAAAGGGAAATA
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AGTAATCTACTTTGCCAAACCCCCACTATTTAGAAGTTTTTAATGTCTTCCAGTTGTGTACTAAATACCTTCAAAAACC
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Fig. 6.29ß

TTCTTAAATACTACTCTTCATGAAAACTTCTCAGATATCCCAGCCTGAGTTAGTCATTCATCCTTTATTTTTGTATCTT
AATGCACTCATATTAAGTGCTTAATAAATGCCTATTTAAGCATATCCAAGGAGAGTTTGCCAAAGATCCCTAATGTATG
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CGTCTGTAAAGTGAGTCTGTTCGACTTGACAATCTCTCATTGTCCTTCCTGAACCAAATAAATATATCTATAAAGAGAC
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Fig. 6.29

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Fig. 6.300

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Fig. 6.301

307/375

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TACGTGTAGCTGTACATACACGTGTATGTATACGTGTAGCTGTACATACACGTGTATGTATACGTGTAGCTGTACATAC
ACGTGTATGTATACGTGTAGCTGTACATACACGTGTATGTATACGTGTAGCTGTACATACACGTGTATGTATACGTGTA
GCTGTACATACACGTGTATGTATACGTGTAGCTGTACATACACATGTATGTATACATACACGTGTATCTATACATACACGTGTATGTATACA
TATATACACGTGTATACTTATATATACATACGTGTACATACATGTATGTCTACATGCATGTATGTGTGTGCAT
ACATGTATGTCTACATACATGTATGTGTGCATACGTGTGTGTACATGCATGTGTGTACATACGTGTGTGTGCATGTATG
TATATGTCATGTATGTATGTTGCATGCATGTATGCATGTGTATATGTATGTATGTGCCTGTATGTATACATGTATG
TATATATGTATATATGTGCCTGTGTATATATACATGCATATATGTATATATGTACATGTGTATATAGATGTATGTAT
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TATGTCCATACTTCTATGAAGATGGGTCTAGAGAGTTGTGAGTAGACGTGGTCCTCCTATAGCTAATCCTTGGGTGTTG
TCAAATAGTCTAGGCTCCCAGAAATATTTTTATATTCATTTTAAAATTAGAACATACTTGTTTCTCTATAATTCTAAAA
GAAGTTGTCAAGGGTGAATTGATAGATGGATGATAGTTTTTAGGCTGAGGAAAGAAAGATACTAAGATTTCAAAATTAT
CAAAATCTTGGACTTTAAAATATATTTCAAAAATATTATTTTAATTTTCTATATGTTTATCAATTTCCTAATCCTATTG
ACAAACTTCAGAGGAGAATATCTTTCTAGTAAGAGTGAGTGAAGCACAGAAGATGAAAAAGTTTGCACAAGATTCCAT
TCGGCCAGCGCGGTGGTGCACACCTGTAATCCTAGGCACCCCATAGACTGAGACAGGAAGATCACTTCAGCCCAGGAGT
TCAAGGCCAGTCTGGGCAACATAGCAAACCTCCATGCCATTGAAAAAAAAAAAAAAAAGGATTAGATTCAAACGTAAGCC
CCAGCCTGTGACTCCTTCACAGTGCTAGTTCATGCATTCATTTGTCATGCTAAAATTTGCAATGTGTGATATTCAAAAG
CAGTGGGTCTTACATGGCCACTAATGCTTTAAATATAGTAGGATTAGGGGTCTTTGCTCCATAAAGTAAAAACTTTCTT
```

Fig. 6.302

CTAATTAACTGTATCTCCTTTTCCTTATGATTTCGTAATAGTAAAAACAGATAAAGATTGCCACCTAGTGTGTTTAAGAT
TTCAACCAATCATGAAATGAACTTCTATAGGAAATTATATTTTCAGGTGCTGATGTTGGTTGAACTTATTTTTTTAAGA
TGACAACAGGCTTTTTCATTTTTAAGTACTTGAACCATATATGGATCTCAACAAAGTCACCATAAGAGTGTGGGCATTT
GGATTTCTTGCTTTTAGTTTCTTGCTTGTTTCTGTTTCCTTATTACAGAAAGAAAAAGGGTAGAAAATAACGTGATAAA
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AAAAAAAAGACAATAGTATACTGTTGAGCCCTGTATTCTCCTGCTTTCAACAGCGCTGGCTCAGGACTCAGCAAGTCTT
TCTCTTTGGGAGAGACAAGGCTCGCCAGGACTAGTGAAGATGACTCTGGTTCACTTCCTGGTAACCAGGGAAGGTGACA
GAGCATGTGGCACCTCCTTGGACATCATGAAATTTAAAACCAATGTGACACTGACTCCCACAGTGACCTACAGTGGCTC
ATCCTGGCTGTTCTAGAAGTTTCTGCACTTGCATTCTGCTGTGACTGTTACCTTGGTACCAATCTCATCTCTCACATAC
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TTTTTTCTGAGCCTTCCTATCCCTCATTCATTCTCCGCCTGTCCTCCATGAACACTCCAGTTCTTTTTTTTAAAAGAGC
TGTATTTTCCTTAAAACAGATCACAGCTGTTTTTTGTTAAGGGTTGTTGTTGTTATAATTGTTTTTATTTAAATGAAAG
ACAGTATTGCATCATCCTCACCCAAAATTCACATTACCAGGCGGTACAATTTCAAAATGTCCAGGCGATATTCACTGTG
TGACTTTGAATCTAGCTTTTCAGTCACTTTATAGAGACCCTAATAGTCTCTTCCCATCTCAGAAGAGACTCAGAAAAAT
ATTTAGCCAAGTAGGCCTAGGAAGCCCCAATTTATTACATTGTATTGTAATCCCTCCCAGCCCCACTTTACGTGGCAAG
GCGCTTTAAAAAGCTTCTGTGGGTAAAGTTGTTCAGGTGCAAGATTTCAGAATTATTTTATTTCATAGGATTTTTTTGA
GCATATGTGATTATTCAGTTCATTAAAAACCCACACATACTTTAAACCAAATCATATTCACAAAGTGAATCTTCTGTTGA
TCTTAACTACCAGATCTGTTTTTAAATGATTATAAGCAAAGTGTAAGACACAAATTCAAAGATATGTTTGTATTATATTT
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TAATATTGAGTTGAGGACTTAGACAAAGATTTTAAAATTATATGTTAGGTATATATTCAAAGATAATTTAAGCATGACA
TACTTTTGAAAATAAATGGATAGGCTATAACCTGCCGTTTATCGAGGACATCTGCCACTGAAATGCTAGGTATCATTTT
ACTGTTACTAAAATGAGACCATCTTGAGATATTTGGCAGAAACAATTTCTGGGCAATTTATTAAAAAGTAATATATTTC
CTTCACTAAAATGATAACTGTCTTTATGCTTTCATAGGGAAAAAAGGCTGGTAAATTATCATAAATATTGTAAAGTGTA
TTACTTATAATACTTATAAAGGAGATACAACCAAAGAGAAAATTAAATGTTTTAATAAATATGCAACCCTTTTGAGTTT
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GCCTTGTCAAGATGACTTAGTCCTGAAAAAAATAAACTTTATAGTAGGTCAATTATAAAGAATAAAATGAAAGAAGAGA
CTATAGCTTGGGACAATAAGAAGATGTCATATTATTTTAATTTTGGTTATCTGAACTGCCTCTCATCCATTTTACAATA
CAATGTCTTCAGCTTGGCCATGCACTGAAGCAGATGAATAGTTTATCAGACTGGCAAATCAAGTCGGCATAGGTGGCAT
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GAAGGGAAAGTCAGAATGATGGGCCAAGGAAGAAATGTAGCTGGCCAACCCTGGCATCACTTTGTAATTTGCATTTATT
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CCAGAGCTTTTTAATGCTGTCATCGTCCCTCTCTCTCTCTCTCAACTACCACTGATTTAGTTTAGGTCCTCATCTCTT
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CCCAAACTTCCATCAGAGAGTAAAATCCAAGTTCTTGTACATTTCATACAAGGGCTCTGTGATCTGCTGCCTGGCTCGT
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AGTTTTTAGACCTTATGCCCTCTCTGAACCTCAGTTGCCTCTTCTATAAAATGATCATAAAACCCAATGTCCTACCTTC
TTAGAGTTCTAGAGTTCTATGAATAATCTCTACAAAGAGCAGGTGCCCCTTGAAAAACATGATTCTGGAACTAAGGGGG
TGTAAAATAACACATGCTTCCTTTACATTGTTTTTTTCCTTCTTCTGCTTTTATTTCCCACATTTTTATTTTCAAGATCT
CCACATGGTTTGATGACGACCACAGCACCTATTCTAGCTATTGGTAACAATTTATTGAGTAACATCATCCATATGCTCC
CTGTGCCCCAGCATGTTTCATGGGTGTTGTTTTGGGTCACTCAGGTTGTTTTCAGATTGTTCATGAGTGAGCGAAAAGT
TTATTTGCACTGTTCTCAGGGCCATGTCCAGGGAGCTGTCTGTGTTGAATAACAGCATTCACTTTTGATTTAAGAATTT

Fig. 6.303

TATCTAGCTAGATCAAGAAATAGCTGATGGTGACTCTGGAAATCCTTGTCTCAAATTATTCATGCTAAGAAATACCACA
AAAGAACCCTGGGCCCCAGTTTCTCACTTTCCTCTAACTGGAATGTTGCACAACTTTCCTTGCCCTTCCCTGTCCCTCC
AAGACACGTCATGCTGACACCTTGCTCCCATACCCCAGGCACGGTGTAGCTGGGGTTGCTGCAGCAACAGCACACAAGG
AGTAACATTTCCCAGTGACTACAGTGGCTGGGATGAGTCAGTTGAACATTTCACTGGGTTTTTATTAGTGTATGATGCA
GTCCCATGTCATCTACAGTTCCTAGCCTGTCCAAGACTTACACGGGATGATGTGGAACACTCCATGCTAATCACCTTCA
AATATTAGATGGTCAAGGGTTCTCTCACTGAAGTCACTGCATTGCTGGTTCCTTCCACAGCACCCCTTACTAATATCTA
CAGTAAGAGGCCAAAAGTAAAACTCAGGCCCCACTGCTGGAACACCATTTCTACTATTGCATATCCTGTGTGCCTAAAA
AGCAACAGTGACCCTGGCTGATGATCAAGAAGCCCCTCTTTTTACCCAGGGAGTTGGACAATTGTTAACATCCCAGAAG
ACATATTTTTAACCCAGTTGCTGGGGAAGAACATGTGCTGAAGTATAAGGATGGAATCACTGTAATTCTCTATGGGAACT
CATGAAAGTGGAACTTGCCGTTTAATATTGAACGACACTGAGCACAGAGTTATTGATGATTTCTAATGCGGGATTTGGT
TTATTTTTCAGACACGGAGATGACTTGATTGTGACTCCATTTGCTCAGGTAAGCACAGCTTGGTGAATGGGCAGGTTTC
TCACAGATGTAAAAATTTAATTTGGGGAATTAGTTCGGGTTATTAATTTAATTTAATTTTAAATCAAGCACAAGTACAA
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GATAACATGATCTAAAATGTTTGGGAATATAAATATGAAAGAGTTGATGACCAGATGTTCTATGTATAGACCAATACTT
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GTCACTGAACTTTTCCAAGCCTGGTCTCTTCCTCTGTTAAATAACAGTTCCAAGAGTACCCACCTCATAAGATTGCTAA
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TCAAGTAATATTTATTAAGCTGCAACAAATAGATTAGTAAATTGGACTTGAACCCTGCCATCAGTGCTCTTAAATTCCA
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TCAAGATATTGGGAAATACACACATATATGTAATTTAGTCCATGTTAGTAAGCTATTTAAATAAAAAAGATGTGTACTT
AATATACTTGAAATTAGAACTTGGAATATAAATCAGCTGTATAATTTATAGGCATATCTGATGCCTAAAAATAATCCCA
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CCGATTTTTTTTAAGGGGCTTCATCATGGTTTTCACAAATGGCCTTTTACTGGAGTATAAGACCTGGAAAGCCCCACTA
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GACCCAATCACTAACTTCTCAAATTTTCTTTAATACAAAGTAAAGCACTTACCACTTAATGGTAAGTGCTATTTTAGCA
TTAACTATTTAAAACTAAAAATATAAGATACTTATGTCACTTGAAGTGATACCAATCTAATTTGTCCTGATATACCATA
ATTGCTTTCACCAAAGGACAGAAAACAATGGATTTTAGAAAAGTCACTCAGAAAAATATTTAGCCAAGTAGGCCAAAGA
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GACAAAACAAAACAAAAACTTTTTGCATAAAAGTTTAGAAAAATAATAAGAAAAGAAAGGAAAATTTTCTAAAATTC
CAAACCCCAAAATAACTGCTACTGACATTTTATTTTATATCCCTTCAGATTTATTTTTCTGTGCATATTTTCTGTCCCA
TGGCTCCTAAAAATGGCCTTCCCCTGTCTAAGGTGTTCAAAGATCCCCTCCCAAATCCATCTCATCTTCTGCCTCTTCT
TGAGAACTCATTTCTTTATGCCTGAGTCCATCTCCAAAACACGAATCATGGTTTCCAAATCATGGTTTCCTTTGACGTG
AAAAGAGCAAAATCCTAAATTAGAGGAGAATTTAGGAAGCCGTGCAAAGATAGATTGCACTTTTTTGGCCAGGCTAAAG
CCTGAAACATCTTGGATAGTAATTTTCAGAAAGGATCCTATAGTATCTCTCCATAGTTTATCCCATTGTCTAAAACACT
GTCAGGGCTAGAAATTCTTGCTGTGGTGTTTAACCTAAACTCTTTTCCACCTCTGATTTCTTATGCTTTCCCCTGTAAGCT
CAGAATCCCTTTGGTCCTCAGAAAAAGTGACAGCTTTAAATATTTTTCTTATTTCAATTGTTAAAGTATTCTCTTGTGT
TTGTGTTGAAATCTTTTGGTGTGGGTGTTTCCTTAGCACGAAGGCTAGAGAGGAATCCCACTGGAGTGCACGCGGCAAC
TGACCTTTTTTTCTTGGCCAGAGTTCTTGGGGGCCAAGGAGCACTAAGGAGGCACAATGCTGATAAACTGTAGGAAACT

Fig. 6.305

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CTTCTCTGCTGTCAGCTGCCTCCGGGCTCAGTGATCCCAGGATTTGTGACACAACTACCCTGTAGTAAAACTCTAAGTT
GACTAGAATTTTCATGTGGAAGTGTGTAATGTTAAATGAAGTAGACTCTGATGAAAGAAAGAGAAAACAGGAAAACCCA
AGCATTATGCTGGATCCCTGGAGAAACCCTCACTTCATTCAGACTAAGGAGATATCCCTGATTGAAAATACAGAGGCG
GGAGAATAACTTGAACCCAGGAGGCAGAGGTTGCAGTGAGCCGAGATCGCGCCATTGCACTCCATCCTGGGCAACAAGA
GCAAAACTCCGTCTCAAAAAAAAAAAAAACAAAAAAAAACAAAAAAAAAAACGGGGTAGGAACAACTACATCTCTTTACCT
TTACTCTAACCAAAATACAGCACCTGGGATACTTCTAGGATAAAAATGCAGTAGTTATTAATACTATAGAATATTACCA
TTTAGTTGGTAATAAAATATAACCAATATTCTCATATTCCTGCCTGTCCCTCAGCCACTTCACCCAGCAGCAACGATCT
TGTGGCCTCATCACCATCCTCAAGCTTGCAGACTGTGCTAGGCCCTCCCTGCCAGGCTATGAAGAAGATAGTCACCTC
GTTTCAGTCGGTGCTAAATTTTTAGCTTTTCTTTAGCCTTAATTAATTTAAATTACAACTACTATCTTCTGGACATCTA
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AACAAAACATAACCTGGAAAGAGGTTTTCTTTTCATAAAAATGACCTTTCTATTTTGTAAATAAGCATAACTTGACTCC
AGCTTTTTCCGGAGGTAAAACCGAAGTAATGAATCTCTCCCAGTGCGGGTCGCCGCTTCCTCCGCCGCCAGTGGCACCT
CAACCTCATGGTCTGTGTTTTACCGCCATCAGTGGTAACAAGTGTATATGGTTATGAAAAACAATCCTCGAAACCATT
TATTTTCCTCTTTTTTTTTAGGCTATGATGTTCAGCTGTTTAAGTGTTAAATGATAACCGTATTTTCCTGCTATTTTCA
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GAAGGGGAGACTTTTTGCTTTATACTCTTTTGTTTAAATTCTGAAGCATGTGAAGATATAAATTATACAAAAATACATA
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GAAAATAAGTTGCCTGCATTTCTATGTCTATTTACCCAGCTGTCAACTGCAATGAACTTAACTATTATGTATTGGAATA
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GAGGTACAAAGAGGTTGCATATCTAGTCCAAGTTCACACATTTACTACATAACAGAGCCAGACAGTCTGGGTTGTTAGG
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TATTGAGTATCTTCTGTGTGCCAAGCATTGTGCATGTACAGGGCGTAGCATTTTACATAAGAAACGCATCCCTGAGAGC
TTCTTACAATGCACAATTCAGATAACCTTAACAAAAGAAAACAATGTCTGCTTACTAGCTAGAGTAGCCATGTGGCAAA
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TACAAACTGCACTAAGTGTTGGGAGATTCTTTAGTGGTATATGAAACTAAGTTGCAAGCAATATCAAGTTTTCTCTTAA
CAAACCATTTTCAAGTGTATTTAAAAATGGCCTTAGCTCAAGAAGTTTCAAGAATCCTTTCTTGTTGTAGGTCTTGGCCA
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GTGTTGTTGCCATGTTGTCTGTGTATACTAAGTCATATGATGTCCTGTTAATTTTCTATAAATACTTCGTGGTGATGGT
TCTAATATCAGAAATGAAGCAGTATGACAAATAAATATGGTGATTCCATCTGTCAGAAATCACCTGGCATGATCAGTCC
TCCGCCCAGTTATTTACACTCAGGGTAACTTTATAGTTTCGGCTTACTTCATAAATTAACCGTGGGTGAATAATCTCAT
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Fig. 6.30<u>5</u>

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CACACCTTAAGATCAGCTGAAAGTCATCCTCTCAAGACAACGATGATTTGGAGAAGCGCCATATCATAGCAAAGATCTT
TGATTCTCCTGCTGTTACTGACTTTCAGATCAGCAGCATTCATGGAGCAATTAATAGAATTGTGGTTTATATGACAACA
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GAAATAAAATATTAAAATGCCAACTTCAAATAATTTTCCAGAAGGAAAAAAAACATTTAAATCATATTAAGAGGATAGT
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CGTTATTCAAGTACTTCTTGTGATAGAAAAAAGTTTTTAAAAATCAATATAAATGTAACAATGCAAGCTATTGGAATGG
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TTCCTCTCCTCATCTACTGGGATAATTGTCATTCATTCTTTTGGTTGCAGCTTAAATAAAAATTTCTCTTGGAGCATTGT
CCCTGATGATACCCAACATAGACACACACCCACAAGCCCTAATCAAAGTGATGTGCCTTTCCATCTCTGTCTGATTGTA
CCTTGTACTTGATTCCTATGTTAGCACGTAGCTCTCTGTGCTCCTGTTGCCTGTTTGTCTGTTGGGATGCTCCTGAACA
CTGTCAGTTCCCAGTGAGTAGGAATGTAGTCGGTTCACATTTGTATTCCCTGTGCCTAGCATAATGCAGAGTCTATCAG
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Fig. 6.305

312/375

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Fig. 6.302/

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Fig. 6.308

314/375

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CTAAAATCATTACAATGATGGCAATGTGTGGAATAAGCATATAAACACATTTTCCCAGATGCAACTTTGAAGGAGACAA
AATCTGAATTCTAGGCAGTTTGTGAAGGAAAAGGAGCAATTTCTGGGAAATTTCTTTTACCCCTCAGCCAACCTCAAGA
CAGTCTGTCTTTCTACAGACCCTCCCTTGCTGTTTCTCTTTACTGGAAACTGTTGAGGGTACTGGAAATAAAGGAGAAG
GCCAATGTATCTAAGACCAGTGGATCAGAGAGGAGCCCAAGAACCTTCATCATTTCCACAGAAAGGGCAGCAGAGGAAA
TTAACCTGGCCTTTTGTCATGCTATTATTTCTCAGTGAGTCTATTAAATTATTGTGGCACACTAAACAGTGTTGTCTGC
ATCTTGTTCTTTGGGGATTCCATACAATTCTCCCCTTGGCTGGCTTATCACCTCCCCTTAGAAAATGGTGTCTCAGCAA
TGATCCTTTGCTTGCAATACAAAAAAACCAACTCTGCTCATTAATGCAAAATTCTTAGTGGAAGAATATAGGAAGAATT
CTGAATTCCCTGAGAATTAGTGGGGCGACAGGTGGTGGTGGGGGTGCTGGAGGACGAGGTTTGAAGGAATCAGTGCATC
TTTACAGTGCTAGAAGGCTACAAATCAAGGGGAAAGTCTCATGGCAGGAGTAGTCGAGTGAGATCCCCACAACTGTTCT
TAGTTTCTTTGTCATTATTTCAAGAGTCAAACAGCAGTGAAGGGATCTAATTTGTCTTGTTTGCGTCCTGGGCCTTGGC
TAAAAGAGAACAGGCAGGTGGCTTCAGTTCTACCAAACTATGTTATAAAATGTGGTAGTAAGAGCTGAATGATGGGACC
```

Fig. 6.30

AGATTTTGTGGGTTCATATTCTAGCACTACGACTTACAGATGCTTGTGCTCAAACAATTCATTTAACCTTCCAGAGCCT
GAAATTCCTCACTTTTGAAGTAAGCACAATAATAATATTTATCTCATAGGGAGTTCATGAAAATTATTTGAGGAGATAT
TTATGAAAGGCTGGGCATGGTTTGCTCACGCCTGTAATTCCAGCATTTKGGGAGGCTGAGGCGGGCAGATCACCTGAGG
TCAGGAGTTTGAGACCAGCCTGGCCAACATGGTGAAACCCTGTCTCTACTAAAAACACAAAAAAATTAGCCAGACATGG
TGTTGCACACCTGTAATCCCAGCTCCTTGGGAGGCTGAGGTAGGAGAATCACTTGAACCTGGGAGGCAGAGGTTGCAGT
AAGCTGAGATCGTGCCACTGCACTCCAGCCTGGATAACAGAGGAAGACTCTGTCTTAGGAAAAAAAAAGATATTTATGA
AAAATGAAGCAAAATGCATGGCTCACAGTAAGCACTTGATCATTGCTGGTTAACTATTATTAAGATTATCTAATAGGGA
TACAAAGTATATTCCTGAAAGAAATTAGGGTGCTCTTAGATAGGGGAAATAAATGCTAGTAGGATTGGTTTCAATGCAC
TATTTATTTTATTAATTTATTTCTCACTGAATTTCTTTTTATGATACTCATTTTTTTTAGAGATCCAGATGGTACCTGT
CATTAATGTATCCCTGAGAACTCAGTTCCTCTGAAGGCACAAAAATAATTTGACCATCTCATCATTTGATCCACACACA
TTTATTGATGCCTCTTTTATGCTTGGTTCTATACAAGACGCCGTGACGTGTGGCCCCTGTAGGAAGTCCTGGCCTCTGT
CTCTAAATCTGTCATCTCAGCCCAACCCCTATTCAGATTATCTGTGCCCTCTGCGTAGCCATTCACTTCGCCTGGGTGC
AATCATGCCTTCCTTACCCAACGTTTAACAAGTCCAGCCTAGCAACCTACATCATGCCATCCAACAGAAGGAAGGCATT
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CCCAGGGGAGCAGTGAGGCATTGTCAGAGGGTGGATTCATTAAATCAATACAGAGAATCCTGGCTGCACTATTTGTTAG
CTATTTGACTTTGGGAAAGTTATTCTCTGAGCCTTGATTTCTCATTAGAAGACGGGAATAATAGCAAACCTATCTCAGA
GCATTAAGTAAGAATTAAATGAAGTGAATTAAGCACGAAACCTAGCATCTGACGCATAGTAGGAACTAAAAAAGAAATA
CTAGTTCTTGAATAAACATTTTTTTCTTGACTGTACTGTGATACTTCCATCAGTGAGTTTCCAAGGCAAATGAATCATC
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AACAGCCCTCCCTCTGGAAGGTTTTGTTCTCCATCTGCCACAAATCAGATTCCTTAGGAAGATATTTGATTTTGAAACA
ATGTTAAAGTACTGTTCATTTTGTTCAAATTTCATTTATGTACCATTTTTTAAAGTGATGTAAATGGACAGCCACAAAA
AGCTCAGCAGCTGGTCAAAACAAAACATCAAAGTCACCATTGAAATGGGGCAAAAAAATTAAAAACTAATATGCTGGGA
CAATGCCAAATAAAAACGATAGTCTGATAAACATTCCTCAGACACATTTTGCTCATAACAACTATTTCCTTTACAGCAG
ATTTAAAAATATGGGAAAATAGGAATATACCCCAGTTGCCACTCTGAATCTTAGCTGTCCTGAGTTCACTGCAATGTATA
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CAAAAAAAAAAAAAAAAAGATTGTTCATTTAAAAATCAATTCGATTGGCCAGGCACGGTGGCTCACACCTGTAATCCCA
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TGGGACTACAGGCACATAGCACTGCACCACCATACCTGGCTAATTTTTAATTTTTTTTTTTTTTTAGAGACAGGATC
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TAAATTTCCATATATAGATCAATTGGATGTTTTGTCCCCAGCTTCCTAGGCCTTTAATAAACTGAATTGTTTTGGTATC
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CAGAATGTAGTCCAATCTGCTTTCAAATCAGACCTTCAAAGAAGGGTAGATTCCATGTGACAGATCCCCTTAGGTG
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GCACCTCTGAAAAGAAGACCAAGAGAAAACTTTAGTATCTCTTTGCTGGAGATGCAAAGCAAGATATAGAAGGAACTGG
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TAAAATTTGAAGCACGCCTGCACCCTCACCTGAATGGACTTCCTCCTTGGCCAGGGCACTTTAAAATTTAACCTGAAAG
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GATAAGAAACATTTAGGATCTCTTTTCTTGGAAGCCTGCTACCTGGAGGCTTCATCTGCCTAATAAACCTTTGGTCTCC
ACAACTTTTATCTTAACCCAGACATTCCTTTCTACTGATAATAACTCTTTCAACCAATTGCTAATCAGAATATGTTGAA
ATCTACCTGTGACCTCGAAGCCCTCCCCCAACTTTGAGTTTTCCCGCTTTCCAGCTTTCCAGATAGAACCAGTGTAAAT
CTTACATGTATTGATTGATGTATTATTTCTTCCTAAAATGTACAAAAACAAGCTGTAGCCTGACCACCTTGGGCACATG
TCTTCAGGACCACCTGAGGCAGTGTCACACGTGCATCCTTAACTTTGGCAAAATACACTTTCTAAACTGATTGAGACCT
GTCTCAGATATTTTGGGCTAACACAATGAATATGAAAAAACTTTTTTGTTGGCGGGGGTGGGAGGTGGGACGGAGTCTC
TCTCTGTTGCCCAGGCTGGAGTGCAATGGCGCAATTCGGCTCACTGCAACCTCCACCTCCTGGATTCAAGTGATTCTCC

Fig. 6.310

TGCCTCAGCCTCCTGAGTAGCTGGGATTACAGGCACACGTCACCATGCCTGGCTAATTTTTGCATTTTTAGTAGAGACA
GGGTTTCACCATGTTGGTCAGGCTGGTCTTGAACTCCTGACCTCATGATCCACCCATCTCGGCCTCCCAAAGTGCTGGG
ATTACAGGGCGTGAGCCACCACACCCGGCCAAAAACAACTATTTTTAAAGAGCATCTAAGCTCAGAAATCACAGGCATA
TTACGCCTGCCACTAAGGGAGTTAGTTCCATTGAAGGATATAATTAAGAGTGAAATGAATGGTGTGCTAAGCACTTAGG
GATAGTGGTTCACAAATTTGTCTGTTCCTTAAAATCACCTTGGGGGGCAGGCGTGGTGCCTCACACCTGTAATCCCAGC
ACTTTAGCAGGTCAAGGCCAGAGGATCACTTGAGGCCAAGAGTTCCAGACTAGCCTGGGCAGCGCAGTGAGATCCTGTT
TCTACAGAAGATAATAATAATCACCTGGAAATTTCTTTTTTTTAATGCAGCTAATGAAGGAAAAAAAAATCCGAGGTTT
TGAAACCAACTCAAAGACAACAGCGATATTCAAGCATAAGATGTAATAAAGGTTGTACACTAGATAGCTAGATAGCCAG
AATAAAAAGGAAGAGATAGTTACAAAAAATATAAGGAGGATAAATGTATAGGATTTCATAACTGCTAATCATATGATTT
TACTGAGTAGGTGTAAAATGCTTCTGATAATGTGTGAAAATTTATAATCCTTCGTATTATATGTAGGATAAACATAGGT
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TGTCCTGTATATTGAATATTCATGAAAATCTCTGAAAGGTGTTATGCTTATTCTTAACCTCTTAAAGGTGTACACTGAA
TGTAATTAAATCATTTTTGCTGGCTCTGGTTCCTCATGAACATCTGCTTTTGTACTTCCCTGTCATTCACAAATGCACT
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CTGCTGTCCCATATTGTCTGGTGGTCAGTTCATGATGTTACTAAGCTTGGCTTTATTGGCATCTTTTGTGAGCTGCTGC
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TTTAGTTTAATGTGCCATAGTCACACTTTCTGTATTGGGAGTGTTAATGGGTGATAACTACTCCAGAGCTTTAGGATTG
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AGCCCATACTTAGCTAAAGTGGGTATTTGTCTATTCCTGTCTGCATACTGTGACTTGGAGATGCCTATTATTTTGCTTG
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CCCAATGGAGAGCCACTCACGCAAATCAATACCCTTTCCTTCTCTCAGTTGGAGCCAGACATCTCTAACTATCCTCTGA
ATTAAACTGGATTTTTAAAATGTTTTTCTGTCCCTAAACTCTTTAAATCCTGAGGTCATCCTTTCTTTGTTCCATCAGT
CTGGCTCCATGATACCAATTACTCCAGCTTGTTAAAGCAGTTATTGGCATATGGTAGTCATGTCTTTTGTTTCTATGCA
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CCTCTTGGTTTGGACATTATWATTTGAAATGAATATTCTTTTTAAATGATTGGAAACTTAGTCGTAAATTCAAGTGGTT
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TGGCTGACCTTGAAAACTACAGCTGTTTAGTCACTTTGAAAACAATGCAATACAAGTGATTACTAGGCTTCAGTTTTA
ACCATTTTATGCTGACTGGTGGAAATTCTAACCCTTCAGAAAGAAAAAAAAATTGTGAAGCAATGGAAATGTACCATCC
TGGGTATTAATGTCATTAATTTTCAATACATTTTCTCACAAAAAGTATAAAGAAGTCTTTGCTTGACCTTACGGGAAAA
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TAGGCATTACATGGTAGAAAGATAGCATTCTGTTCCAAGAAATCTCTATTCGTTCTCTAGTTGCTGTGTATATATTCTC
TTTTAGCTAGAATATGGTTAGTAAATGAGTCCTGAAAATTTCGCTGTATATACCAAAACATGTTGTATACCATAAATAT
ATAAAATATTTTTATTTCAATTTTTTAAAATTGTGCTTCAATCAAGACAGAAGTTGATTTCTTTTTCCAGAAATTCAGA
GTGAGCAGTCCAGGGCTGCTGTGGCTGCTCTAAAATCACCAAGGGCCCAGAGCTTCTTTCAGATAATTTTTTTTTAACA
TTTACTAGATTCATAGGGCTTTTGTCTTCATGGTCTGAGTGACTACAAGAGCCCTAGCCATCACAACTAAGTGGCCAG
CAGCAAGAAGGAGGAAAAAAGAAGGAATAGTACACCCACTCCTTTTAGAAAAGGCTTCTGAGAAATCCCACATAATGCT
TCTACTTACATCTCCTTGGCTCAAATTTAGTGTCTGCCACAAAAGAGAATAAGACACACACACACACACACACACAC
ACACACACACACACACACATATATATATATTTTTWTTTTTTCTGGCAAGTCTCTAGTTCAGGGATTTCAATCTCCTTTA

Fig. 6.31;

AGCAAATGAGTCTTC   GCTAACCTCTCCCTCTGGCCTGCCCAGTCTCTCTCGAATCTCATTGCCATCCTGTCCTCCTT
ACTCACTTTGGGAAA CCAGCTCTGTCAGTTTTACAGTCTGCCTTTTCCACTTCCTTACCTGA TTCCTGGGAAAAAT
AATCCCACATCACTTCAAGCTGGAATTCCAACAAAGTCAAGACCCTGTCACCTCATCGGACTCTCAGAGCACTTGACAGT
TAGTTCCTGCTGTGCCCCTCTCTCAGTCCTCTTCATGCTAATTCCAAACCTGTCCTCTCTCTTAAGCCTCCCAAACCAG
CAGCTCTTCATCTCCTTTCAGCAAATAACTTGTTGCTTTTTGAAGAAAGTCTGTGTCAAGGCATGAACTTCCTCAGCTC
CTACATCCTCTTGCCATAGATGTTATCCATAGGTCTCTTACTGACATCCATAATCCCTTCCTCCCTTCTGGTCTCAGCG
GATGACGTTTCCCTTCTCCCTCCTGCTCTAAGTCAACCCCTGCATCTGGGCCCTTGATCCTCTCTCCTCCCAGCTCCCA
CAAGTCCTTGCTTCGTCTGTTTTCCTTTCTATCTCTCACATCTTCAATCTTTACTTTTATTCTGGCTCTTTCCCCATTC
ATGTCTCTCACAATCTTAAAAACAATACACAGCAAACCACAGCAAAGTTCAGCCAACTCTTTATTGATCCTTTAGCTAG
TTCTTGATCTTCTCTTTTCAGTCAAGTTGCTCTACCTCCTCATCCCTATTCAGTCTGCAGTCCACTGTAGTACGACTTT
CTCACAACACCTCTGTTCCAGCTGCCCTCCCTTGGTTGCCAATGAACTCCTAATTGTCTTCTTAGTTCTTAGAGGCTTC
TTCTTCTTAAACTCTTTCTTTGTTCATTTACTGAGTAAATATTTATTCAACATGTTTTTGTTGGGAGCCATTTTGTTCC
AGATGTTTCCACACTCTTCTTGCTGCTCTGAATACTTCTTTTCACTATTCTTCATGGTCTCTCTTCCTCCCCACTTTAC
CTCTTAAATGTGAGTATTCTAGAATGTTCTTTTTGATTAATTGGT TCTCACTCTGTGTACTCTGTAGATGTTTTCAT
GTACCTCATAGTTGTATTTTCCATCTTATGAAAGTGGGTCCCAAAACTCTGTCTCTACTCAGAGTAGGCACTTCCCCAA
TTTCCAAATTTATTCACACAAAAGCCCAATGTCCATTGATATCCAACTCTTCAAATCCAATATGTCCAGAGCTGGACTC
ACATATTTCTTTGCAAATCTGTTCTTCCTATCTCAGTTAGTGAAACACTTTTAGCCAAGATCGAAACTTATAAATCATG
AATCCTTCTACCAAATCTATCTGTCTGTCTATATATATTTTTGAAACAGAGTTGTCTCACTCTGTTGCCCAGGATGGAG
TGCAGTGGCAAGATTTAAGCTCACTGTAACCTCCACCTCCTGGGTTCAAGCAATTCTTGTGCCTCAGCCTCCCAAGTAG
CTGGTATTACAGGTGTACACCATCACACCCAGCTAATTTTTGTATTTTTAGTAGAGACAGCGTTTCACTATGTTGGCCA
GGCTGGTCTCGAATTCTTGGCCTCAAGTGATCCTCCTGCCTTGGCCTCCAAAGTGCTGAGGTTACAGGTGTGAGCCACC
ATGCTCGGCCCCAAATGAATACAATTTTTATTTTATTTTATAACTATTGATCAAATCCATGCTCTCTTCTCTAACCTGC
TAATATTTCCCTAGTTCAAGCACTCACCAACTCTCCCCTGAATTATAGTACTAGTACCTGAAATAGTGTTTTTGAAACT
GTCAGTTGCATTTAGCATTTTATTTTAAAAATATGATACAGAATTGCGTAAAAAATAAGAGTATAAAACCACATCTATA
ATATCAAAGATAAATATGGTTTTATAAAATTATTGATACACCCACAGACATGCAAACACACAGACATTTTGCCATGA
AAATGTATTTCTTACTATAATTTTTAAAATTAGAGAACTACTTTCTTAAATGGTCGATCTCTGTTGCACCTCTTCTTAG
CTCTGGCCTTCCATGATGCCAGCAAATTTATGTAGTTGAAAAATTAATCTGCAAATGTCACTCTTCTTAAAACCTTTTC
AAGCCTTCCCATCATCCATGATAAAATCAGATCTCTTTCGTGTGGTACCCAAAGCCCCTATCCAGATATGTATTGTACC
TCCTCTCCAGCCCAGCCCCTGGCTGACTCTTTCAGCAGCAAACTTTTTCCACCCACGTGCACCATGTTTCACAACTGTC
TTCATTTACTTGTACTCTTCTCACCATCTAAAACTCGCTTCTCTCTCATAATTGCCTGGTTCACTTTATCTCATTTTAG
AGGACACCTAAAATTCAAGCCCAGTCCCCTTCCTCAGGCCGAGTCAGGCTCTTCTCCCAAACCTACCTCTAGCATAACC
TAATGAAATTGCTCTAAAAATTATGGTCTATGCCTCTTTATTAAAAAAAAAACAACAACAACAAAAAAAAAAACCCTGCC
CACCAGTATAGCAATTCTTTTGTTATAACCACACATAGTACTAGTGTGTGGTTCTGGGGCACTGTCATGCCAACTTACC
TGTGTTCAATAGACCTGAAGTAGGGAAGATCAGCTGACTCTTTTACATCTCTTCTCAGAACATCTTGAAATGATTCTTG
GGTGCAATATAAGAGCAAAACATGGAACAGTAGAAACCAGCATGGAAAACTTTTTCTCCTATTTGAGAAAGTGGGATGA
AAATGTGTCTCTTAATCCCTTTGGCAGAAGGGAGTAACTTGAAAGAAGTGGATGATGGATGATCTTCTCCACTCTGTCA
GCCCTCTGAGCAGAATGAGGAAAATGTGAGAAGCGATACCATTGCTATATTTGTTTCATTGACCATTATTTTAAATAAC
TTACATGGTCCTGAACAAGGACATGTTTTTGTGAAAAGGACAACATTTCTTTTGGTGTGTCCGAGTAGCATTTAAGAAA
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CATGGAAACACTAGGGAGTCATGCCTGGATCTGACAACTGTGCAGCACCATGCCAGGTGCACAGAGAGTTGTTGTGAT
GAATGTCGAGAGAGGTGGTGGGGGATTTTTGTTGTTTTTGCTCCTTCCATTTTCTTTCCCTCTGACTTGGGCTGTGGTT
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CTTTCCTTTTCTTGAGAGGAGAAGAACAACTCCACAGTGCTTTCCTTTTATGTTTTGGGCTAATTTTGTTGTGTTGTTT
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TTCTTTTTCCAGTCTCTTCCCTCACTCCTTCCCTTATTTGTCTTGTTTCCCATTCTTTGTCTAAATCTCCTCTTTTCTC
TCCCTCTTCACCCTATTCATTCCCAGCTCCCTAAAATTATATCATTCTTCCTTCATCATGCTCCCTCATTCCTCACTGC
AGCATCACCCAGTCCAACTCTGTCCCATCTGTGATTCCTCACTTCTGCTTTCGCACCATCTCTCTCCCTAAGGTAGAAG
TGGTGCCCAGCTTTAAACACAAGGCAAAGTGAGGGTATGTTCCACTAGGACAGCGTGTGTCACTCTGCTGCCATACACA
GGGAGTACCATCAGTGATTATAAAGAGGAATTTCATTCCTGCCAGAGAGCTCACAAATTAGGTTATTGTTGATGTTCAT

Fig. 6.312

ATAGTTTTTAGTGACTGGAAAAAGATAATGCCTCCTTCTATGGTCACTACCAGGAAGTGAGCTCCATGCAGCAGACAGA
AACCAACCAGATAACTGTGAGATCCCAGCCAAGCACAATTTCTACATGTTAGTCAATTGCAAGAAACACTGAGATGTAA
TGTAGTACTTTGGACTCATATTACTCAGGTGACTATGTTTTCAAGCAGCCAGTACAGGACAGAAGAGATTCATTTCCTT
GAATCAAAGCTATTTGGTGACTTTTCCTGGAAACACTCTGAGGCTTACAGGAAGGGATTCCAGATTCCCAAGCTACTGG
CACAGAGTAAGCATCTTTTCTTTACATAGTAATTCACAAACGTCCCTCATCACCATATGACAATATCCTCACTGGATCA
GCTCGGTTACCAGAAATAACTAGATCAAAATAAATGTCATTCTCACATGGACACAGGTGGACTAGGGTTTAGAAGTTTT
GGGAACCATCACAGTTCTAGGCACCTGAAACTTCTTCTTTTGCTTGTAGTAAATATGGTGTCTTTCTCAGGACTCTGCC
CTTTGTTTATTTATATCTGGTCTGGTTTTTGTTTTCTAGGCCCAATTTTAGGCCATCAGTAGATTTAAAAATTTTTTTTT
CTCTTGACACCAATTACCTTAGTTGCAATTTCCCATTTATAATTGCAACTTGAGCCTACAAATTTACAAATACAATGAG
GTAACTTTCCACTGTGGTTAAATATTATAGATTATTGGCTGCTGGTAAACCCATGAATTGCCTGCCTTTGGATTAGATA
TTCAGACCAGGTCCATTTAACAATTCAGACATTTTTTTAAACAATACTTTTTTGAGAGTCTTCCATGTGCCAGGCCCTA
ATCAAAGCTCTTGAGATACATTGATGACCAAAAAGAAGCATTATCCCCATTCTTACTCCTTATAATCTAGGGTCCAGCG
ACASGGCGAGATATGCACTGTATTAATGAAATAATCTGTACTGAATGTAAAAATTCCAAACTGAAGCATTTTATATGAAG
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TCTTCCTCTTGTTCTTCCACCCAGAAAAACCATCCTTATCCTCATATTCCTCCTCCACCTTCACCACCACTACTACCTA
TTTTGAGGCTGCCACCACCACACTATAACATATTCCTCCCCGCTCTAAGGCTCTGAATTTGGACCTCCCACATGACAGG
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TTGTGTAGCCATGGCTGGGCTAATTTCTGCATTTGCATTTTAAACACCCTGAATCAGGATTACACCTGAAGACTTTCAA
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Fig. 6.313

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Fig. 6.316

320/375

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Fig. 6.315

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AACATGATCATTTCTACTTGTGGGTTTGGACTGACAAGGACATTTCTAGTATATCTTGTTGACATCAGGTTAATAAGCT
CTAAGCAAAGCTGAAAATGAATGCTACTTCCCACAATCAAGTGGAAATGTTAATATTATCAACATGTCTTAAAGGCCAT
GAAGTCCTATGCCTGTATATGTAACAGGAATAGCAAACAGGAACTTTTCTTAAAGGGAAGATTAAACTAGTGTTATTTT

Fig. 6.316

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TATGTATTCGTTAATTGCATGCTCCCCATTTTTAAAACTGTACCTGAAAATATACTGTAATAAAAAATTAAATTTGAAA
TATTATACCATTGAATAAGGTCCTCTGGATAGTTTTGAGATTTAATATTCTTTATTTCAATTTACATGTATACTGAATC
ACGTAAAAGTTATTTAGGTCAGTACTTCCCCACATTTGTTCCTTCATGACATACGCTGATAACTATATTTCTACGAAAA
CTGCAGTAAATAAAGGATGTGGCTTGTGTTCAGTCTCCCACCCACTAGCGACAAGAGCCAAGGGAAATCAATAGCTTGG
CTCACCTGTCATCTCCACTCTTTGGCACACCATTGAGAAGTCTAATATGTGCAAAGAAAAGTTAATTTTACTTTCAAGG
GTTATACCTTTGCTTATTCCAAAGAACGGTGGGGGAGTGTACCAGGGCCATTATTATAATGTATATTATTAAACTTTTT
ACATTTTAACTGATTTCTCTTATAGTAAAATTAATTATATATTAGCTTTGGGGTATACAAAATTGCACTTTGATAATAA
TATTTAGGAATTGGCAACTTTGTGGGCAAAGTTGTGGGAACTATCAGAAAAGTAAAAAATAAAGGTGGGACCCGAAAGA
AGGAAGGATTGTAGGTTGGAGCCATACGCGACTAGAAACTAAGACATGTATCCTTAAAAACCCTCAAATCTGTGTTCTG
ATGGGAAAACCCTGCTAAGTCAATGGTTCCCAGACTTTGGAATTTCAAAACAAAAAACTGTGAGGCCCAACATACAATT
ACCAACTCTTCATTTTTGCCAAACAAAACTATTTTCAACAGAACACACTAACATATACTGGAACCATGATTTCATATTA
TGACATTTATTTTCCACATTCTATATACTTTTAATATGCAAAACAAGGACAAAAAGGGCAGTCATCTCCCATGGGGGGC
TTAGAAATCTCTCATTAATGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTTTTAAATTGGCC
TTATTATTTCTTTAAATTTAACCAAGGGCCAGTAAAACTCCCTTTTGGAATACCACACGCCCGTGCTGTTCAGGAATGA
TTATGATAAAGTCATTCATGAGGACATAGAGTTATTTCATATGCTCTCAAAAATGACTTGCAGTGCGAAACACAGCTAC
TATTGGACAAGACAACAACCCAACTTTATATAACAATTTTTTAATTCACTAGTACTTGTGTATCAGTCAATAAAGTACC
TGGTACTTGTGTACTAGGCACTGGGATGGAGTTGGGGATACGAAAGTGAAAAATATAGTCACAATTTCTACCTTATACT
CTAGTAGGGGAAAGAAAAACAAAATGTATGATTGAACTGCCTTAAGCACCATGAAGGCAACATACAAATTGCTGTGAGA
GCATATAACCAACTCAGAAATCAGCCTTAAACTGAGACCTAAAAAGGACTCCCCCTGTGCCTTTTGTATATTTAAGGCA
TCTTTTTTATATTCTAGAAATACAATATCATTGAGCAGATAGATTCCAAAGGGCCATCTTCTAGCTGGATGGAAATAGAA
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TGGTTTTAAAGAAGGGAATAACATGACCAAATTTGCATTTAACTTTACTCAATAGTCTGGAGTATTTAAATATGATTAA
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AAAATTTTTACTAGGTAGGTAAGGGCAGGATGATTCCATTAGAAACAAGAACTTTTTCTCCAGTGCTCTGGGAAGAATT
GATTGGTTCAGGGCTAAATCACCAAGATGAATACCTTCTTTTGATTCACACACCCTTTTGACATCTGGGTTTCCCTGAA
TCATATGCAATCTAGGACTTACGCAGACCACTCCAGTTTCACTTACGAAATTTGGGTTAGCCAACTTAAACAGTCATGG
CTGTAATTTGCTTCTGAGTTAAAAATGCTATTATTATTTTATGTTATTTATGGATTCCTGCAAAGGAGAAAGCATTTTTA
TTTTGATAAATTTTGTTATTAATGAAATTAACTCAATTTTGTAGTAACATAGTATACAGTGAAAATGTGCTTTACGAGA
TGCCGTGGGCTTGTCTCTCAAATGGACAATGAGGCTTTCCGAAACCTTTGTGCCTTTAAGTTTAAGGAATTGATTTGGC
GTCTTATCTCAGTGTACACAAAGCTTAGTTATCACCTAGGAAATATTGGATGTTTGTCTCAGGGAGAAGAGACAATTTT
TGAGACCACAATTAGGTTTTGTAGCTATTTGACATCCAATTATAATCAGATACTCAGTGAAACATGCATTCATAATGCA
GGCCAATGATCATATATTTGGCTTAGCTTTTCACTAGAAAAATTCTTTAAAGAAATATTGTCCTTCCAATTTTAGTGGT
ACTGAGGAACAGCTCTGGTTATAAGGGGGGAAATACTATATGTCCTTTGTTCAATCTTACTAATCTTTATTAATTTTAT
GATTTTCCCCATTTATTTTTTAAAAATCATTAATAAAATTATCCTTGATTGAGTTTCTCCTATGTGCTAGGCATTGTGA
TAGGACTTTAGTAAACATCATCTCTAACCCTTTCAATAACCCTGCAAGGTAGGGTCATATTTTATAGATAAGGAAACTG
AGGCTCAGAGAAGTTATGTCTTGAAATTATATAGTAGCAAGTGGTTAAGCCAGGTTTCCAACACAGGTTTGTCTGACTC
TAAAACCAGCATCTGCCCCCATCTCCTCACCATTCGCGACCAGAATGTGATGCACGTCTACCAAAAAGAGAGTAGCAAA
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ACCATGGTGAGAGCTATGTAGTAGATGAAGCAGAATGATGGCTGGGCTCAGGGAGCTCATACATTTGGAACCCAGATAA
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TGCTGTCTCTACAAAAAATAACAAAGAACAAAACCATCTGAATTTCATGCATTCCAACATATATTATTTTGTTCCATTC
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TTGAGGTTTGGTCATTAGGTTTTGAATAATCAGGAGCTATGTGTTGTAGTAGGAAAAGCATACCATTTAGAGAGAGACA
GCTTTAGTTTGTCAGTCCTGCCTCTTAATAGGTGGATGACCTTGAGTAAAGCATGACGCAGCCTTTGGGCATCAGTTTC
CTCACTTAAAGGTAGAAGGGTTGAACCAGGTGTAAGCTGCTGTTCTTCCATGTTATGATTCTAAATAGTTCTAGATGGA
ATTATGTTGCACATTTCATATGTTGTTAAACAGCTTCTCTCTGGAACAGTTGCTCCTAATCTCCTAATAAGCTTACTAT
GGGTCACTCAGCAATCTTCAAGAGAAACAAACATTATTGGTTGGTGGATACCAAGACAATTCTGATGTGTTTGTGTCA
AAGATTGGTCAGATGAAATCTAACTAGACATGTATCATTCCATCATCAAGCGAAATGACCTTGACTCTAAACATCTTAA
TTTGAACTGTAGACATTTCCCAAATGCTTTAGGGTAGATTTGGACCATAATACACTCCACTGTAAATGGAGAAAGTTCA
TCATCCCATAATAATTGCCCCCAAACTGCTAGGTAGGAATGCATACTGGGTTTAGTATGTATTAACTCCCTTGTGAAT
TCAGCTTTTGAGAGCAGACCTCTTAACAAGTCATGAGTATTTAGAGTGAAATATTTAATTCTTAGTGATACTATTTTCC
TATTCTGCTATTATTATTCAGCTATGATTATTTTCCTTCTCCTCCCAACCTCCTATTTCTTTTTCTTTATTAAAAAAAG
GAGAGGAGGTGCATGCTGCCATCTGGGGTGAATATGGGCTTCACGTGGGGACCTGGGCGCTGGTTCTGATCCAGAGAGT
GAGCTAGGAGCACTTACACAAGAGGGTCTTAGCCACTGTGTCGGGGGAATAGCCTTATATCAAAGACTGAAGCCTTTTA
GTAAAACCTGAAGAATAATCAGAAGCAGCATTGACTAGTCTCTAACCTTGGTTGAGGGACACACACACACACACACACA
CACACACACACACACGGCTCTGAAAGCTCAGACACGGAGGACACGCTGTGATCCACTTAGCCCTCTCTTCTCCTTCA
GGGGGTCCCTTGCAGGACAGAACGCTAAGGAACAAAGTCTTAACTGGCAAGCTGCCGGCAGAAGGAGACTGTCTGGTGG
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Fig. 6.317

ATTGCAGCGCTGATTAGGAACGGGGAGTCCTTTGCGTTTCAGAATCCGGTCCACATTTTGCTGAGAGCCCAGGCACTGT
GAATAAGAGGAGCCTCTGAGGACTGGTTTCTGAT‍TCGCACTAGGCATGATAAGAATGCCATTGACGTCAGCAGAGTCA
TCATCAGGGAGAATGGCACCCCAGCTCTTGAAACCACACCAAGCAATTGCAGTGCTTTTGAAAATTAAAAGCCGGTGCT
GGAGTGTGGGGGGTGAGGTGGGGGGTGGGGGAGGGAAAGAAGCAAAGAGGGAAGTCGTGGCTCCAGCTGAAC
TCTTCTTTCTTCTATGCGTTTCCTGTATTTTAAAAAGGCAGGCTCACGGCCCTCCTGTGCACACGCCCGCTCACTCACA
CACACTGCTGCCTGGGCTCATTAACCTAAATTGGGGGAGGTGGGGGAAGAAGAAAGAGAGAAAGAAGGAAAAAATTAA
TTCTGTTAACACTTCTGCATGCACATAAAACAAACTGTGCCACCCAGGTGCTGGACAGAGGCCCTGAGCCACATAAGCC
TCTGAGAATCAAATTGCTGAGGTAAAACTGATATGCTTTAAATGTGTGTTGATGAAGAGATTACAAGGGAGGAATTAAA
CACTTCAAAGGAGCTACTACAAAGCAACAATCACTCTTCCCCCATCCCTCACACACGCACATTACTGTGGACGCTTGCA
TGTTTGTGTGTCTGCCTGAGAAAATGGCAAAGTTGGTTTAGGAAGGGAACACTGCAGGCATGTTGTGATTTCAGCTTTC
ACAGGCAGCTCCAAGAACATCACCTCCAGCTATAATTCAACGGTTACCTGGACTGGAAAATGTCTTGGAGGTTACGAAG
TCTTTGAATTCTTTAGTGTAAATTGTCTCCGCTTTGAAATTCCACAATTGATCGAATCCTTTCATTCTCTTTCCCACTC
CTAATTGTTCCACTCTAACTCTCCACCCACCGCCAGGAATATGAAAACAACATAGAGTAGAACTGATGATATCAGCTGT
ATAACACGCTTTATTTTACGGCATTAATGGGCAGCTTTCCAAGTCCATAAAAATCTTTTGCTGATCTGATTTTAATATC
CATATTCTTCAGTTATACTAGAAAGAATGATTTCCGAGATGTGGAAATGTTAGCCGGGAAACAAAAGGACATAGGCACA
TAAAGCAAAATCTGGCAGGAAATGTGGGTTTAATAAAATTGAATTATGTGAAACTCCTATTTCAAATGGACTGATACTAG
AGGAGCAGCCCTCGAGAGATATGCACAGTAATGATTCTTCGTTTTCAAAGACCTGTTTCTTTTTGAAAAGGAAGAAGCC
CTGAACAGCATTCCTTTCAAATACACAAAGGTAGCTAAAGTACAACCTTCTCCTCTTTCCTATTTGTTCTTTAAAAAAT
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CTTGCTTTCTCTCTCTCTCTCTCTCTCTTTTTTTTTTTTTACAATTGTACTTAAAAGAATAAAGGATTCTGAAAATC
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TGTATTATTGTATTTGGGGGATCAATAAGCAGAGCCAGAGCCTTCACACAAGGCTCCTTGTGCCCCCTCCCCCGCTCCA
TCCCATGAGGTTAAAAAGCACTGACTTTGACTTTAACACTGTGAACAAAATACAGAACAACTGTTTCAATTCCCTTCAC
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AACGAAGAAAGACTTTCTCCCTTCTTCCCACCTACACATTCCTTTTGTCCCCCAAGATATCCCTTCCTCTTGCCTGGAA
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AACAACCTCTGCCATAAATTTAGGAAATAGTCAACCTGTTCACATTTTCATCTTTTTACTATATATGTTCTTATTCACA
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GGCGAAATGCAGTGAGGTAGAAGTGAAAGCAGCTTAGAGTCAGCCTGGGCTAAGATAAATGAGGAGTGGAGCTGCAGCT
GGAGGAGCAAGAGAATATGGTCTACTCGAGTATAAATCAAGCTTGCTGGTGAGCAGTTTAGGACTAGCTCCTGGTTTTA
CAGCAAGACTTTTTATGTGATCTTGACCAAACCACTGCACTTGATTGCTCCTTTCTACTCCTTGGTCCTCACGTGTCAA
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AATGAAGACCTAGTAAAACATTATAACTTACAGATACCTGTCAGGGCAGAACCAGAAGCTGGACTTGGGCTAGGAAAAG
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TTGTAGAAATATTAGGAAGTGGTATAAAGCCAAGACCATGTTTACTAAACAGTGCTGCTCTTTTTTTTACATCATACAG
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AACAAACCACTTCCAGTCTCTGACATGCAAGCCCAGAAGTTTTCAGATTTACAATGGCATACTTTTTTTCTATCTGCTA
AGTGTTAGATGTCACGGTCGTCTGCAATGGAGTAGAGCTTTAGAAATGAGGATAAAACATTTGGTTTGATGTTAGTATTA
AGAACTCCAATATCCAAGGCAGGAAACTTGGTAATTAATATTTTAAATGGACTGGAGGGGTCACAGGTATTAAAATCAA
TATTTTGATGGCAAGATAACAATGTAGGGACAAATAATTGGAGATGTTTAGCAAAATTATGAATTAAGTAGATGTATAA
ACGGCAGATTATGAAAGGGAAACAAAGATTACCTGGGAAGAGTTTTGTGTTTTAGTCAACTGAATTACCAGAATGGGCA
GAAAAAGGAAGTATCAAAGAAGGGTCTGCTATAGAAAATTAAATAAAGAAGAAGATGCCATGGAAACGTTTTTAGTGC
AAATACACTATTGCTCATGGCTGAAGACAGAAAGAAGACTATGTGTGACTTCCAGATGAAGTTCAGGCATAGAGGCAGG
GCTAAATAAACAAGCACATAGACCCGAGAATTCTCGTTGATTGATCTTTGGAAATTCGGTATAAATATGATCAGTTCAA

Fig. 6.318

CCATGCCTTAGCTACTGGATCAAAGGAGACATCTACCGGCTTTGGGAGATAGGAATCTGGAAAATTGACAGACTGAAAA
AACCAAAACCAAAGAAAAAAAAAAACAAAAAAAACAGAGAATGCAGAATGCCATTTTAAAAAC¨¨AAGAGGCTGTGTC
TGATGTTACAGGACTCTTTGACTCCATTAGTCTCAATTCTAATATTTAAGTTAAGTGATTGTGGGCATAGAATTTTTTT
CAAAACCTGTTTCGAAATGAAGGCTTCCCATACATTGAAGTTAAAAGTGGTCTTGAGAAAAGAACTTCTTAATAGGTAA
GGGGTTTTACTTTGGAATGTTGAAAATGTTTGGAACTAGATAGAGGTGGTGGTTGCACAACATTGTAAATGTGCTAAAT
GCCACTGAATTCTTGACTTTACTATTTTTAAGTTTATGTGATATGAATTTCACTTCAGTAAATTATCTTTGTTGGTGCA
TAATTACTGTATATATTTATGGGGTTCATGTGATGTTTTGTTACATACACAATATGTATAATGATGGAATCGGGGTATT
TACGGTATCCATCACCTCAAGTATTTATCATGCCTGTGTGTCCTCTTTCTAGCTATTTGAAATATGCAATACCTGTTG
TTAACTCTAGTCACCTAATCTGCTATGGAACATTAGAGCTTATTCTATCTGTGTGTTTGCACCCATTAATCAATCTCTT
CATCCCCTCCCCTCCCACCTACACACTCCTCCCAGGCTCTGGTATCTATCGGTCTACCTTCATGAGTTCAAGTTTTTTA
GCCCCTGCATATGTGAGAACATGTGATATTTATCTTTCTGTGCCCAGTTTATTTCACTTAACATAAAGATCTCCAGTTC
CATCCATGTTGCTGTAAATGACGATTTTATTCTTTTTTATGGCTGAATCATATTCCATTGTGTATACCATGTTTTCTTT
CTTTCTTCCTTTCTTTCTTGCTTTTTTTTTTTTTTTTTTTTTTGAGATAGAGTCTTGCTCTGTCATCCAGGCTGGA
GTGCAATGGTGCCATCTCAGCTCACTGCAACCTCCGCCTCCCGGGTTCAAGCAATTCTTCTGCCTCAGCCTCCCAAGTA
GCTGGGACTACAGGCGAGTGCCACCATGCCCGGCTAATTTTTGTATTTTAAGAAGAGATGGGGTTTCACCATATTGGCC
AGGCTGGTCTGGATCTCCTGACCACGTGATCCGCCCGTCTCGGCCTCCAAAAGTGCTGGGATTGCAGGCATAAGCCACT
GTGCCTGGCCGTATACCACATTTTCTTTATCCATTCATCTGTTGAGGGACCCTTAGGTTGATTCTCCATCTTTGCTATT
CTGTATAGTGCTGCAATAAACTTGGGGATGTAGGTATCTCTTTGATACACCAATTTCTTTTCCTTTGGATAAATACCCA
GTAGTGAGATTGCCGGACATAGTGGTTAGTTCTATTTTTAGTTTTTTGAGAAATCTCCATGCTGGTTTTCATAGTGGCT
GTACTAATTTACATTCCCACTGGTGATATGTAAGAGTTCCCTTTTCTCTGCATCCTAGCCAGCATCTGTTATTTTTTTT
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GCATTTCCTTGATTATTAGTGATGTTGAGCATTTTTTCATATTGGACATTTGTATGTCTTCTTTTGAGAAATGCCTATT
TATGTCCTTTGTCCACTTTTAATGGGATTGTGTGCTTTTTTACTATTGAGACGTTTGAGTTCCTAATATATTCTGAATA
TTATTCCCTTGTCAGATGAGTAGTTTGCAAATATTTTCTCCCATTTAATAGATTGTCTCTTCACTCTCTTGATTGTTTG
CTTTGCTGTGCTATACAGAAATTTTTAGTTCAATACAGCCCCATTTATCTATTTTTGGTCGTGTGGCCTGTGCTTTTGA
GGTTTTAGCCATAAAATCTTTGCCTAGACCAATGTCCTGAAGTGTTTCTTCTGTGTTTTCTTCTAGCAGTTTTATGTTT
AAGTCTTTAATCCATCTTGAGTTGATTTTTGTATGGTGAGAGATAGGGGTCTACTTTCATTCTTCTGCTTATGGGTGTC
CAGTTTTTCCAGCATCATTTATTGAAGAGATCTCCTTTCCCTAATGTATATTCTTGGTGCCTTTATAGAAAATCAAATG
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CAATTTTTTTAAAAATGGTCTTGGCTACCATAAACTGTGATGTTCTTGAGAGCTTACATTGTGTCTTGTTGATTTTGGA
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ATTTTGGAGTATTACAACCTCATTTGGTTCTCAGAAGCTTTTTTTAATTTCATCTTCTCCACGCCAGTTTATACAATTC
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CTCTCATGCCTTACTTTACTTTTATATTTTTTCCTCCATTCTTCCTTGTTTCATTCATTCACTCTCCCAAGTCCTCTGG
CCTAAGTTTTATTTGTGTGGTATTAAGCAGAGACCAACAGAACAACTGTGAAGACTTCTTGGGGATAAACCAGGTTTCA
CTTGAATTGTGGTGCGTCTGCCATTGGTGTGGAAGCAGGGCAGAAAGAGGACAGAGGCATACCAGCAAGTACAGACATC
TAAACACAAAGAGAGGAGAACAACCTCAGGGATACTCCCTCAGTGCCTGGGAAGAAAGCAAAACTTAAAGTGCTCATAG
AATTGTAAATAAACTCTCTCCTCTTTGAAATTTATGTTCGATGGTTGCTGTGTAGTTGCAACAGTGTCATAAAGTCCAT
GAGGCCTATTTTTTTTGAGTAATCAATATTGTGGAAAAGTGCTAGAGCAGACCTACCTTATATAAATTAATTTGTTTGA
ATTGCTCTATTCACTATGAATAAATATTTATTGGAGTACCTTCTATATGCTGGGGACCTCAGCCATGAGATACGCAGGT

Fig. 6.319

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GGAAAAAATAACACTTAAGCAGCTTATTGTATAGCAGAGAGGTGGTTGAATTGATAGACAATTAAAATACAACGTGTTG
AAAGGCCTGTTAAAGCCACAGGGTCACTCAGTGGCAGGAGGGATTCAGTCAGCCTGGGCCAGCCATGGCAAGTAAGTGG
CCTTTGGAAAGTTACCAAGGTAAGACATACCTGATACGTTCAAGAAACAATTCCAAGGTTCAATAAGGCCAGAAGTTAA
AGGGCAATGTAAGCCCTGGAAACTGAGGCTAGAAAGGTAGAAGGATCTCTTTTTACCACACACTGTGCCGGGCACCATG
AAGAATACAAAATAGCTAGCTATTATGGAGCCTACACTTTAATGTGTGTACATAGATACATGTCCATGTTGGAATTTAA
CTCCTTGAGAGGTCATGAATTACTTAGTCCCGGAAGCATTTGAATTTATACTGGAAAGCTATAGTAGAGAGTTGACAAA
ATATTTTAAATGTAGTAAGAAGAATTTGACAATCTATTGTGTGGTATGCTAGCAAGCCAGTTCTCCTTCCTCCTTTTCT
CACCATCCCACCACCCCTCACAAAAWAAAAAAAAAAAGTTAAAAACCTGATCTGCAGTGTTTGTCTATGGCCATGGTGC
AAATACTCCTCCCATGGCTGGTTCCAAGCTACCACTGGTTCAACAGTTCTTTTGCAGAATTCTTGAATCTTTAACACTT
AACAAGATCCAGTTCCAGCATATTCCTGCAACAGCGATTCTGCAGTTCTCAATGTCAAATAATAGAACCAAAAAGTCAT
ATAAGTACTACAAAGAAGAAGAGAGGAGTTGAGAGCCAGCTTAGCCAGGAAATGCTCCAGGGGGAAGTAAGGCTTGAGTT
GGGACTTATAGATAAGAAATGTAAATAAAAAGGAACAAACACTCCCACCGTGATGCACAAATATTGGATCATGTATGTG
TGTGACAAGTGTGGCCGTCTGTAGTGGAGGGTCCAGGTAAAATTTAGGCCATGAAAGCCTTGGTCTGTCAAACTAATAC
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GCCTCATGTAAGTGAGAGTAATTCAATTCAGCAGTAATCTTGGAAATTTGTAACTTCTGTGGTGAGAAATTGTATCCTG
GGATGTTGGGTAAAGCCACTGTGAAGCCAAAGAAATTAGTGGCTAAACGTAGAGAAATTAAGATGAAGGAACTGGCACA
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TTACCAGCTGTTGTTAGTTATTTGAAATTAGGTGCCTAAAGCAACCTCTCATCTTGCAGAAAGTCATCTTTCTTGAAAC
TTTTTAAAAACTTGCTTGAAACATGGAGACTTGGAATGGGACGTCTATCATAGTAGCACATCTGAAATCCTTCTCATTC
CTGCTGTCATTTCTGTCCTTTCGCAGTCCACTTTGTCACCACCCCCACCATTCCTTGTTCTGCCTGCAATCCAGCTTTC
CAGTCACTTTCTCTACTTCTTGCCTGGCTGCAGCGTCATCATTTAAACAGCTCTGGGTTGGTATCTCCTTTCACCAGAA
TTCCTGCAGCCGTGGAGTACGTGCAGAGCATCTTCCTTTTGACCCTCTTAAAATGCTGAGGGGTGTTTCCGACACACAT
TGCTTTGTAATTCAATAATAGGGCAGTTGTGGGTTCTTTTTATTGCAGTTATGCTGATTTTTAAAAGCACTCTGAGTAA
GAAAGGAAAGGTAGTAAAAATATACGTGTACCTCGATGCTCATTGATCACGTTTGCACATGTTCTTGCCAAATGTTGTTT
TTAAAACCTGTCTCTGCAGGTCTGAGCTGTGAAAGTTTGGGAAATCAGGGTCAGAAGGCAAAAAAAATTCGTTTGTATC
ATCGTCATACAATATAAATTACATGTAGCATATACATAGAAAACAATCCAAGGGATGAGCTTCCTGGAAATCATTCCTT
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GTACGTTTACACATTTTTTGGCTGTATATTATATTTACTATATATATTTATTAGGACTACAGCAAACGCTAATTCATCG
ACTGAGATAAGAAATGTGTTTTCCATTAATAATTATAAATCTGCCAGAAGATTGTGAGATTTACATTGTTGGGGATCTA
AACAACCAAAATCATTGCAAAGTTATTTTAAAACTATTATTAATTATTTATGTAATCATAGATTATGGCTTGCAGAAGA
CTGAAGAAATTGTTCAACAAACCATGTAGGTTTTGTGGAAGAGCAGTAGACTTTGATAAGTAAGCTCAGAGTGGTCCAT
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CAAAGGAGGTCTCTAAGCACTCAAAATAGATGTTCACAATGTTCACTCACTGAAGATAATGTAGGTCATTAGCCAGAGA
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CCCAAATCATCAAAAAATACAAAATTATATTTTGCTCAATCTGTATTGCTCTTCTATATTATGGTAAGATTTCTGTTTA
GTGCACTTAAAAGTAAGACTACCCCCAAATTTAATCCCCATCCCCTTCAATGTCTCCTCCTGGACATTGCTCCATCTAT
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AATTCCCTTCCTCTTTGATCCTGTATCTTCCATCAGTTATTACCACATATCTCTATTGATTTTCGCACCCAAACTTCTA
GAAAGTTTACACTTATCATTTTCTTTTTCTTTTCCCCCCCACTTATTCCTCTACCCCCTTGAAGTTAGCTTGCAGCCCCA
CCACTCTGTTGCAACTACTGCTGTAGTTCCTCTGCTCTCCCAATTGCCAAACCAAGTAGATATTTTTCAGTTCATTTCT
CATCAAGTTTCTCTAAGTCAATTGCCAGTGTATCCCTCTCAAAAACCTTTCCATCTTTGCATGAGGTCGTACACTCTTT
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Fig. 6.32C

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<div align="center">Fig. 6.32[ ]</div>

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CAGTAATCACACATGTAAGGGCTTTGGGTAAGATGGACCTTGGTTATCCAACGCTTACTGTGTGACTTTGGTAAATTA
CCTAAACTTTTCTAGCCTTGGATTTTAATGTGTAAAAGATAGATAATACTAACTCAGTAATTCTTTATGGGTTCCATGA
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```

Fig. 6.32Z

CAATGTAAGACTATTGAGGTTTAATTTTGATTGTTTTTTCAAAAGTTTGTAATCTGTTAGGAACAATTTCTAAAGTAAA
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TGTAAACATGTACATGACTTGTGGTGAATATGGTGATTCTCACTTTATAACCAAAGAGGGTGGATGTTACAGCATATGA

Fig. 6.323

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GAACCTAGGTCCCTGATAAACATGGTGTCACCATTCCAACCTTGACCTGTTTCTAGATTATTTTCTTTACTTTTCTTTT
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TATGAAAACATAATCACCATATGTGAAGAAAAAACCACCAATGTTTAGTACAAAAATTGGGAGGGGGAATATTATATTA
GAGAAAGTTTATTAATCCAGAAAACCACAGAATTTAAAAAATATTGGAAGTTGGGAATTTGGGAGTTAAAGGTACATTT
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CAAACAGGCAAAGGCTAAAAAGGAACACGTTAGTGCTCAAAGTTTGCTTCTTGTGCATGCTGAATGGGAAGAAAAATAA
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TATATTTAGTATCCAAGTTGGGGATAAAAGCTAGTTTTTAAGATTTTCTGTTCAGAAATCTTTTCTATATACTATTCAG
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Fig. 6.324

330/375

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TCCTTTAACTGTGCTTATCCCGTAAGTAATTGCTAATGTTCTTAAACTAATCGAGAAAATCATTTCTATTAGTCCCTAA
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Fig. 6.325

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GCTTGACRTGGTCAGTGGGTGTCTGAACATTTAGTGCTAGTGTTTCCTCTCATCTCTCTGCTTTTATTTGGACTTCTCA
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TTTATAATCTGCTTTTAAAAATTTCACAGTATATTGCATCATGAACACTTTTCTTACCATTAAATACAATTCTCTATATA
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Fig. 6.32G

ACATTTTAATGGCATTTCCTTACATGAATATACTACACTTAAGCAATTTTCTATTATTAAATATTTGAAGCTCTCCCCA
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TTGTACAGCATATTTGGTTTTCTTGTTTGGTTGCTTACGTTAGAATTTTCTGAGCAACATTCAGTAATAGTGGAGACAG
CTTCTTTTTGTTTCTGATTTCATTGTTAATTATAGATGTCGAAGAAAAGCTATTCATTACCATGAAAAGGAACTATTCT
ATTCTTATTGTACTGAGAGTGCTATTTTACTATCAACTCTTTAATAAGTGTGATATACTACTGTATTAGGCCATTCTCA
TGCTGCTGTGAAGAAATACCCTAGACTGGGTAATTTATAAAGAAAAGAGGTTTAATTGACTCACAGTTCTGCATGGCTG
GGGAGACCTCAGGAAACTTATAATCATGGCAGAATGCACCTCTTTACAGGGCAGCAGGAGAGAGTAAGCACCAGCAGGG
AAAATTCCAGAAGCTTATATAACCATCAGATCTTGTGAAACTCATTCATTATCATGAGAACAGCATGGGGGAAACTGCC
CTATGATTCAATTATCTTCACCCAGTCCCTCCCGCAACACATGGGGATTATGGGGATTACAATTCAAGATGAGATTTTG
GTTAGGGCATGGCCAAACCATATAATCTACATTAAGGACATTGTTAAGTGGCAGGATAAATGGTGTGGGCTCAACTTTA
TACTATAGAGGACCATTATAGCAATATCATAAAAGTAATACTTTGAGAGACTAAGCCAATAATAGGACAAATATAAATT
TGAGGAAAAGATATACAAAACCATTTAGTAATTCAAGCATCTAGTGATAATTGTATAACCATATGGACAAAGGTAGAGA
AAAATACCTATAGAAAGTTTGAACTTAAGAGATAGGTCAGGTAACAAACCCACTTTTTTTTCTTTTTTTTTTTTTACCTGA
TTACTTCCTGGTCATGAGGTTCAAAATATTTGATAGCAAAATGATTTAAAGTCCAAATTAATATTGAATTGAATATACA
CATTACAGCCAAAACCAGACCTATTTTTCTGTTGCTTATCATATATAATTATAGAAAGCATAACTGAACAACAGGAAGT
TATTACTTCTTAAATATCAGAAAACATTTGACTTTCTCTTCTGAAATAAGGAACGGCAGTAACCGAATTCTAGTTACTG
CTCTTAATAGTTTTGTAGTTCTCATAAAATATTGCTTATATATTATTTGGCTCCCACACTATCCATTTTGATGGTTTTA
AATTTTTCAAATAAAAATAAATGAGTTTTCATACTATGGATGACTTGACATCAGTTAAAAGTGTAAATTCATACTTGAA
GATAGCAGGACTCTATGCTCTTTCCTCATTGTTCTGGTCAAATACATTTCCAATCATGATAATCAATAATGTGACTATT
AATTTTTACTTGAAGTTAATCTATTCATGTTTGATTAGAAGTGTCTAAAACATAAACTTGGGATGTCAATGAAGAACTT
TTTAAAATTCTCCTCCACAATGGCTGATAATCAGGCCATGATTATATTTTGAAGATTTTACCTAGTTAGTCATATTTGG
GGTAAGAATATATGCTATCTCCTTCATATTAATAGTTCAGATCTTTAATTATAACATGTTTCATGGTCTATAAGATTAA
ATTGAACATTAGGAAAGCTTAGCATTTCTCATTTGTGAGTCTTACATTCACACCTGTCAGGATTGGTACAGATTCAAGC
TTGTTTTTAATAAACTTCAAAGAAGATCAAAACTAGGGATCATCTGTTTCATGTCTTTGACGTGATAGAAGTTTGGGGC
CCCCGTTTTGCCCTGTAGCTTTTGCCTAAGGAGAAAGCCACCCTTTTCAGCTGATACTTGCTCCCACAGGTTTTTCACA
GTTTTTGAGTAGCGAGTGGCAAGAAGCAATATGACGATGTTGGTAGGTAGAGAGATACTTCTTCACCATCTGGTGAGCA

Fig. 6.327

TTCTGGAGAACAGAGACAAGTCCCACACTGTTAAAGGAGTGAAGTCATTTTGCAGCCAGTTTTTAAGGATTGATTTAAGA
ACTCTAGGAGCAGGCCATTATGTAGTCATCCATTCAGCACGATTCTTCGTGGGACTGTTTACACTAGAAGGAACTGTAG
TGATCAATTTAGCGTTGAGGAAACTGAGGGCTACAGAGTGCCAGGGTCACGTTCATCACACTTATTTGTGTTTCAAAGC
AGAATCCACACTCCAAAATGTGCAACTACTGTCTTACTATGGAATTTACACTGACACATTCTGAGGTCATGCAGGAAAT
AGACCAGAGTAAAGCATTAGCAATTAGGTCAGTAGCTTGAGATAATGCAGGCTCTGAGGCAATCTTCACTTTCACTGAC
CACAGACTGGGATTGGGAACATGGGAAACGTGTGTGTGTACAGACACAGCAACCCTTTTCTATGTGTCATGTCTTCAGC
TGAGTTTTAAGTTGTACTAGTGGAAGCCATAGCCTATGTTGGCTCATAGGTCCTAGTGCCCTTTCCCCTTCCCTTGCCT
CTGCCCCTGCCCCTTTCCTTGCCTTGCCCCTGCCCCTGCCCATTCCCCTTCTCCTCTCCTCTCCTCTCCTCTCC
TCTCCCCTCCTCTCCCCTCCTTTCCTTTCCTTTCCTTGACACTGGGTAGAGTGTAATGCAATCATAGCTTACTGCAGCC
TTGAACTTCCAGGGCTCAACCAGTCCTCCTGCCTCAGCCTCTGGGTAGCAAGGACAATAGGCACACACCATTATGCCCA
GCTTATTATTATTATTATTATTATTAATTTTGTAGAGATGGGTCCTGCTATGTTGCCCAGGCTGGTCTCAAACTCC
TGGCCTCAAGCAATTCTCTGGCCTCAGTCATGCAAAGTGTAGGGATTACAGGCATGAGCCATCTCTTCCAGCTCTAGTG
CTATCTCTTTATAGGAATCAAAAATTTGTTTTGGCTATTTCAAATTTTGTAGAGTAAGAGGTCAATTAGAAAAGACTCA
GTTATGTTTTCAAAAATGCACTGTGTGCATATGTGTGTAGAAATGTATTAGAATGAAATAACATTTTTTTACAGTGGCT
ACCATTGGTTTGTGAAATTATGCATGCTTTTTCTTAATTTTCTCTAATTTTTCTCTATTTCCTAAGATTTCTACCATGA
GATTTTTCTCTTAATGTCTAGTTTTATTACATAGGTAATAGATAAATATATTCATGTCATAAAATATTTTTAAAATTCA
GAATATGAGAAAGTCCTTTTTCATGCTCCTTAGTCTTGCTGTCTTTCTTGAAAGTATCCACTTAGTGATGGATACAATA
TGGTGTATATCTGCCTTAATCCATTTTGTGCTGCTCTAACAGAATACTTGAGACTAGGTAATTTATCATGAGCTGAAAT
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CCACATGGCAGAAAGCAAAAGGGAGAGAGATAGACAGATTGGGAGAGAGAGAGGAAAGGGGACCAAACGTCTCACTCTT
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GGCATGGTGGCATGTGCCTGTAGTTCCAGCTACTCTGGAAGCTGAGGTGGGAGAATCACCTGAGCTTGGGAAGCTGAGG
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CCTTTTTTAATATTGCAAACCATACTACACTAAACATTGTCGTGCATACTTCCTTGTGTATTACATTGGTTTTTAAATC
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TGGTTTTTGTCTTTGTTGTTCTTATGCAAGAAAGACTAGCTCTTTTTATCTAGAGCTGGAAGGTTGCTGTCTTGGAGTG
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GACTGATATGCATGGTAGTCCTAAGATGTGTTTGAGAAAAGAAAATATTAGGAGCTCTTGACAAATCTTGAAAATCAAT
ACAAAAATTAGCCAGGCGTGGTGGCGTGTGCCTGCAGTCCCAGCTACTCAGGGGGCTGATGCAGAGAATCCCTAGAACC
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GCAGTGAGAAAAATAGTTTTTTGCTTTGATTTGTGTCAGTTATACTATGTTGTTGGAGCTTTTGAAGTAGGGAGTGATCA
GGTAGAGACAGGCAGAAGAGATTAGTTTGTATGTGTTATCCTGAAATCTGAGTTATTTGACATTTTTAAGGCAAGGCAT
TGATTGAGTTTTATGAAGATAAATTGGCAAAGAAAAGTAATCCCATCTGCTCAATAACAATTCTTTGCTTTTAGCCAAG
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TGTTTCTCTCTTTGTCACTGAAGACTTCACGTAGTATAATAGCCAGTATTTTCTGTGATTATATAGCAATATTCTCAAA
CCTAATTGTTCATACAAAGTACAAATCAGGGACTTTTTCTACTAGGTTCTGAAGTACTTGTTCTCTAGATTTAAACTCC
AGTGTTGTACTAATGGGAGTAAGTTTTTCTTTTATCTGCCAAGGAGCTCCTCTTACATGTAAAAAACTGTTGTTTTTCC
TCTTTGTAAATATCAGTGTAAAGTACACACTGTATAAAATGTAAATTACTGATGTGAGAGTGGCCATTTTATTCACATT
GTTCAATGCCAAAGTGGGCCAAAGGATTCTGGCTCATTAACTTTAAGGAAACTATAGTATTCTTTTGTCTCTTTAGCCT
TCCTTCTGGTTTATTCAAAGAATTCTCATCAGGTTGATCAAGAGTTGAAAAGTAAAGCACTGTTCTTAAAAACTGCTCT
CTAGAAAGATCTGCAATGGTTTTGAGGACTGCCAAGCACAGGAGGTAGAAAAATGGATAACTAAATAACCTCATTTAC
AGAAAATATGTTAGGTTAAATTGCATAGGAAAAGATAGTATGGGTCTTTAAAAAAAATGGATATTTATAACAAAGAAAG
GATAAAATTAATAAGTAAGTTATAAGGAATATTTAGAAAAATAAAAACAGTATTCTAAAAGACAATACTTATATCTTTC

Fig. 6.328

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TAATGTAAATCTTAGAAAATGGTGAGAATTTCTTGTATGCTTTATATTTATAAGAAGTAATAATAGCTTTTGAAATAAT
TCCATATTATAAGAAGTCTTTTCTTCATATGTACCCAATTCCAAATAAGTGGATGCTGGAAGAAAATCTGTAATTTATT
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AGCCAACAGCAAATTAGAATTGAAACTCCATTATGTCTCTCAGGTCTTCTGATCAGCTCTTATTATACTTACTCTTAGT
AAAATAATCTCTATTAATAGTGAGAAGTAAGGAACTATATGATATAATAAAAAGTGCTCTAAAGGAAGTCAGAACATTG
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CTGGCCTAGACGCTTTCTAAAATTCTTTTCAGTTTCAACCTTCTATGGCTTTTGGATTCTTTTATTTCTTATACTTTCA
TATTACTGAATCTCATAACTCACTTTTTCCTCCATTTTATATTTTCAACTTTTCTAGTCTAGGGCTCTGCACAAGTTAT
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CTCTGCCTTCAGAAAACCATCAATGGATGTTTATAGGYATAAATGACAAGGACAAAGAAAGTGGGAAGGGAGATAACAG
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CTGGACAGGAGATTGATATAACCTTTGGAGAATTCACCAACCCAAGAGAAAAAACCCAAAGATAATGGTGCAAGGTATT
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TCTGATTTTTAGTACTCTAGCCTTAAATATGAGTAGACAGATAAGATTTATCAAATACTTAGAAAAAGTAAGTGAAAGA
AACCAAAACAAATGGAAAAGACCAACTTCAAAGTACAGATGTTGGCCGGGCGCGGTGGCTCACGCCTGTAATCCCAAC
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TCTACTAAAAATATAAAAAAATTAGCTGGGCGTGGTAGTAGGCAACTGTAATCCCAGCTACTCAGGAGGCTGAGGCAGG
AGAATTGCTTGAACCCAGGAGATGGAGGTTGCAGTGAGCCAACATGGTCCCACTGCACTCCAGCCTGGGTGACAGAGTG
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TCAATAGAAGGGTTGAAAGGTAAATTGAGGACATCTCTATGGAAATGGAGCAAAAATACAAAAAGGTAGAAAATACAGA
AGAAAAGTGGGAAATCAGAGGACCAGTTCAGGAAATCTAACATCTAAAATCAGAAATTCTAGAAAAAAGAGAAAACTG
CAGGAGGACAAAATTGAAGAAAAATTTCCAGAACTCATTAATTGGAGCATTTAGCACAATATAAAGTATTAAAAAAGAG
ACTTGTGACTTGTTCTAAGGATCATTATCGTATCGTATAGAAGGATCAAAAATCAAAAGGGCATCAGACTTCTGAAGAT
TCGTTTTGAAACCTAGAAAAGAATGAAGTGATGCCTTTTAAATTCCGAAGGAAAGTGATTCTCAACCAAGAATTCTATA
CTCAACCAAACTATTGATCAAGAGTGAGCATGGAATAAAGATTATTTCTGAGGTGCGAGCCTTTAAAAAATGTATCTCT

Fig. 6.329

GATGCATACTGACTCAGGAAGCTACTGGAAAACGTGCTCACCAAAACAAGGGAATAAACTCTAAAAACATTACCCTTGG
GATAATAAGGAAAGAACATCCCAAAGTGACAGCT^AGCAAGAGACATGGAGAATAACCAATCCAGGTCAAAGAGGCCTC
TGGATGAGATTTCTTCAAGAAGATGAATTTAATAAAATTCTTGATGTGTTTGAGCCATACTTAGATTTTTGTAATATGG
GAAAAGTTTGGGATTGAATTAGTGATAAGTATATATGGACATCTAAGGGAACAAAGAAACTAACAAAAGACAAGAATTT
TCAAGAAGGAAAACAAAGAAAAAAAGGTAATCAGGGTATGTTACATAGTTTAGCTGCTTATAGTTTTTCTTTGGTTCTG
CTCATGGAAACACAATGACTATCAATCTAAGTAAGACTATAATATATTAGAAGGATGGGTGATGAGAAGTGTGAAGTGT
TGCAAAGGTAAATCCTTATCTTCCGCTATGAAGTATCAATAAGCAATGCCCAAAAAATGAACTATTAAGAAGTAACTG
TAAAGTTATATCATTTAGAGATAGAGTGGAGTATAGCAAATGAATCAGCTAAAATATTTGAAAATGGGTACCCTCTGGG
GAGTGGAAGATACATGTGTATTGTGGGTGGGGGATGCACTGCAGGAGATCTCTTTTTTTAATCCTTGTGGTACTACT
TAGTTCTCTAAACTATTTGCATCTATAACTTTGCTAAAAATAACATTTAAATTTAAAAATTGATCACTCTTGTAATAAG
TTCAAATTGAAACAAGGAGATAACATAGTTGCTAAGTTTTATTTTTGGACATATTTATAACATTGTATATGTGTTAGTG
AGAATACCATGTAACATCACTCTCAAGCAGTACTTCTAAAAGTAGAAATTGCTGTAATATTTCTCAAAAACTATCTGGC
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AAGAAAAGTTATTTGTTGCAACCACAGTAGACCAGGTTAATGGTGCCAAGATGGAATGCGGATAAAGGCTGACAAGGC
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GTTTTCCTGGTTGCTCAGGAATCACCCAGGGAAGGAGAGGGCTTCCCCACGGGATTAGGGAAAAAAAAAAAAAGCTTC
TTGTTCCATAAGAGAGTCTACAGATAGGATGAGTAATAAGGGATAGATTTTACTAAGGTAGAACAAAATGTTAGGACGC
TGGTACGAGCACCACTGAAATATCCCTATATCAAGTTTTAGTCTTTTCATTGCATCTTCTGAACCTGCTGGAGATGCTT
TCACATGGAACGTATTTGCTATAAACTTTTCTTTATCTTTTGTTCAATGCTGTGAAGTTTGCTAATCTTAATGAACCAA
GTCTCTTCATGCTGACAACTCATTGTAAAAGAGGTAAAACTGTGTTTCCATGGTATGGGGAATGGAGAGGTATAAGGAG
GAAGATGGATTTAAATTGATTTTTGGAATGCTTGCTTTATTTTATCAGTTAAAGAAAAGGTCTAACGGATTATTTAGAT
AACTTTAGGCTCCAACTGCACTGATCTGTTTGTCATTCCTTTAAGAAACCATTTTGATTCTGACTACTTGAATTTGCAC
ATTTTGTTCCTTCTCTCTGGAATACCCTAACAAATTCTTTCTGCCTGATAAACTACTTACTCTTCCAGGTCAAGTTGAA
ATGTTCCCTTCAGTGTGAAAACTTTCATGACTCAAAAAATAACAATTATAATAATTACAATTTGATGAGCTACCTGGCT
GTTTACCTAATTCTCACACCATTAGATGAGCGAGGTAGTTTTAGGCTAACTTTAAAAAGGAGGAAACTGAGACTTACAA
GACTTGGGTATGTGGCCCAAGAGTACATAGCTAGATTTGAAACCCAAGTCCAAACCCAGGACTTCCTGCTTAAAGCCCC
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AGTCTTAGATGGTTTTTCATAAATTTTGAGTCATGAAAAACTGACAACATATGAGTCTCCAAGTACCTTTTAATATATG
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CCATGTCCTTTTTCTTTTTTGTTTTCTTTTAAATGAAAACTCATGAGAAATAAGAGGGCAGAATGCATTAAATTATTTC
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CAATATTTAAATGCCATCATATAGATTAGAAATGGAATAAGGTATAGGTAACTTACTGCATTTCAAAAAAAGTACTAAT
TGAAACATATTGTCAAACATATATCTTTTTCTCTCTTTAGAAACCTATGGCTATTTTCTGTCTTTCAGCKCACGCTACA
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GACTATAGTATCTGCTTTTTGCCACAGAATGAAGAATAAACCACGTCTTATAAATAAAAATCTTTCCATGGTCCCCCAC
CACATAAGGACATGACCCAACTTCCTTTGCTAGTCATCCATCTCTCATAATCCTCTCCAGCCCATCTCTGCAGCCTCATC
TCCACCCACACATGCATTCTGCTGTAACAAATGGTGGGCAGGCTTCCAAATGTACTGTGTTCCCATGAAGTTGCACCTT
TGCTGATGTTATTCTCTTTGCCTGGGTCCCCTCCCCATCTATTCACTCCTTTCTGCACTTCACCCTTGCTCTTCTTGTG
CCCTAATAGCTCCTCCTTTTGTGACACATCAAAAATTGCCCACTGTAGGAAGCCCCTCCAAGACTAAGAGTGCCTCTCT

Fig. 6.330

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CTGTGTTCCTGCATGTGCTTTTGGAGCCCTTATCACACAGTTGCTCTAGAATTGTTAGCTTGACCACACTGTGAGTTTT
CTTAAGGGCAGGGACTTATTCCTCTTTGCATATCTGGAATCTTACCCAGTGTTTAGTACATAC::CATGTTCAATAAAT
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GAACAAAACAAATTGGTATAGAATGGACTTTCTTATAAGAGAGTCCCAAATCATAGATCATAAGGACAATCTTATGTTG
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AACTTTAGATTTAATATGTTTATTCTGCAAGATGTTTGGAGACAGGTATACACCCTAAAACCACCATTACAATCAATGC
CATAAACTTATTCATTGCCTCCAAAAGTTTCCTCCCATCTTTTTGGGTTTTTTTGTTTTGTTTTCCTTTCATGGTAAGG
ACACTTAATATAACAGCTACCCTTTTACCAAATTTCTAAGTATACAATATAGTATTGTTGATTGTATACTTATGTTATA
CAGCAGCTTTCCAGAACTTATTTTGCAAGACTAAAACTTTGTAACCCTTGACCACATCTACCAATTCCCATCTTCCCCC
AGCCTCTGGCAACTACCATTCTACTTTCTGCTTCCATCAGTTTGACTATTTTGGATTATACATGTAAGTGAGATCATGC
AGTATTTGTCCTTCTGTGTCTGGTTTATTCCACTTAGCATGATGTCCTCCAGTTCCATCCAACTTGTTGTAAATGACAG
GATTTCCTTCCTTTTTAGGGCTGAATAATATTTCATTGTATATGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTG
TGTGTGTGTGTGTGTGTGTATATATATCACATTTCCTTTGTCCATTCATCTGTTGATGGACATTTAGGTTGCCTC
TGTATCTTAGCTATTACAAATATGCTGCAGTGAAAATATCTCTTTAAGATCCAGATTTCAGTTCTTTTGGATATATACC
AAGAAGTGGGACTGCTGGATCAAGTTACTTCCTGTTTCGAAAGTAGGGCACCCTCTGGACATTTCTACAACTGAAGTGA
TTGATTCTTACTTTTGGAAGTCAGGCTATTTGTTGGAATGGTGGGGAAAAAAGCTAGAGCTTAAAAATAAATCTAAAGT
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GCTTTTTTAGGAGGCTATGTAGTCTAGCTAGCACAGAGTGACCCAGTGAGAGCTTGGTTGGCTAGGACTTTCTGGTGGT
AGCCCACAGAGCTCACCTGAAAGAGTACCAAGGATGAAAATATCATCTTGGCTAATTGGTCTGCTAGTTGATTTAAAAA
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TCCCTCCTTCCCTCCCTCCTTCTCTCTTTCCTTTCCTGGTAAGTATACTAGAAAAAAGTATATTTTTTTCTCATTTTTT
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Fig. 6. 33l

AGGCTGACTCACTGACAATGTACACATAGAGGTGGGCCTCGAGTGTGTGCAAATAACAAGTTTACTACTGCAGGCAAAC
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Fig. 6.332

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CATAAAAAAAGAATGAGTTTATGTCCTTTGCAGGGACGTGAATGAAGCTGGAAGCCATCATTCTCAGCAAACTAACACAG
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CCCCACAGCCATTAAAACCCACAGTCTTCCTTGAAGACCTACATCCAAGTTCACCTTCTTCCTGAAACCCAGCACACTGAG
CTATAGTCCATACCTCCCTCTCCCTGCTCGAACTACAGTAGCACTCCCAGGCTGAATGATTTATATGCTACCTTACACT
GTCTCTAGGCAGCAGCCGATCATGTGTCTGATTTTCTCAGCAGGTTATTATGGTTCCTTCCTGGGCCCAGGCTGCTGAA
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Fig. 6.33̄3̄

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TCTTTCATGTTCTTGTATTTCACAGCAACTATCACATCAGAGAGGCTCAATAAATATGTATGGATTTTTTGACTATTCT
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CCCTGAAGAACAGTTGAGCTAAACTCACAGAATTCCAGGATCATGGTATTGGATGGGATCCTTGACAAGTAACTGGTCA
GTTTAGGAATTCCCTCTACAAACATAGCTGACCCCATCCAGGCTAACTATAATGAACAACCTTAAAACCAACACCTAAC
CCAGATAGTACAATTTCAGAAATCAGATAATAAATGAATGTTAGAGCTGGGTAGGTGCTAAAGATAATCTGTTCAAACT
CCTTGTTTTCTCTAAGGAATACTGTATGCAGTAATTGACAAAGGTGAAGAACAAAAGACTGTTATATCCTAAGATTGAT
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ACGTAAATAGCATACATAATTGTAACTTTTTTTAAACCTGTTTGCAAAAATTGTCCTGTAACTTTGTTCAAACAACTTC
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CTGAGATCAGGGTGTCAGCATAGTCAGGTTCTGGTGAGGGCATCTTCAGGGTTGCAGACTGCCCACATAGCATGTATCC
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CTAATTACTTCCGAAGGGCTTTCCCTCCAAATACCGTGACACCGGGGATTAGATTTCAGCATATGAATATTGGAGAGAC
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CTGCAGCTTCTCCTTCCCCTTCAGGGACTGCTTAGGATTTTCTCCTTATCTGGTCACCAACCTCTTAACTGGTCTTTAC
TTTTCTGATCTGCCTTTATACTATTCCCAGAGTAATCTTTTTGAAAAGCAAATCTGACCATGTGACTTCCTCACTTAGA
AGATTTTTAATGGCCTCTTTAGAAGAAAATATTGTCTTAGAATAGTATACAAGTCCTCCAAGACTAGTTCTCCATCTTT
ATCTTTGTCCACTGTCTGATCTAAACTTTATATCCCTACCTGTTTGTAGCTTCCTGAATGGCCTGTGTTCTCTAAGTGA
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Fig. 6.334

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ACTTCTTCAGAATGAATCCCTGTTCATTCCTTGAAATTCTTTTTTCTTTCTAACTACCCTAAAAACTGTTCCAGGTAWC
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AATTAAACTTAGCTCAAACTCCAATTTTTTGGACAAAAGGCCTCCTTATTCTTATAAAAGCTTTCTCCTCTTTCTTTGA
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GTTCAAAGGTGCCTTGAAGAAAACTGATGCTGTTGTAGCAGGGGCCTTCAAGGCTCAATTTATTTCTCAGTTGCTGTTC
CATAGCCCTCTCCCAGCCTTTAGGGAGTTTAGCCAATAGTTTTTGCKTGGGTGTTTTTCTCTGTTTAGTTCCTTGACTT
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Fig. 6.335

AGCAAGATTTGGTGCAGTTAAAGAGTTTTCATGTATTTTTAAGGAGAGCCACAGTGTATACTACTCTGGCAGGGGTTGA
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CAAATGATCCTGAAAAATCGTGGCTAGCTATATTGCCTACTCACCTAGGAATTTGGAAAAAAGCAATATTCTCAGCTCT
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GCCAAAACTTTCCTGAATTATTTTCCACATTAAAAAATAACAACTGAAATATAATGTGTGGAGCCACATCCTGTTAGAT
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CGGACCCGTGAAGCGCTTGAGAACTGAGTCCCCCTTTCCCTGTCTCTTCGCAGAGGAGGCCTACCAGAAACTGGCCAGC
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CCTTTATATATTCAGTTGGCCATACATATCCGCAGGAAACATTAACAGAGTTAGAAGGTTCCTTATGATCATTACTTTT
GTTTCCATTAATCTTGGAAAGAAAGCCCAGTTTTTTGAGGTCATTTAGTACAAGGAGGCTTCAACTAGGCATCTGTGCC
ATATGTGCTAAGGTGCTGGTCTTGGCAGTTAATGGAGTTTTGAGGGCTGAGAAGTAACCTTCAGCCTGGGACAGCCTTA
AAATAGCACGGAAGGCAAGGTATAGTGTTTTCAGTTTTGCTGTATTTTTTGGATTTCCTGTTCTTTTTGTCTTTAGTTA
TGCAAGGAGATAATTGGAATTGTGTATGTTTTTAGTGGCTATTTGCCTGTGCATGTCGTATGCAGATGGCTGAAGGATT
CAGCCAGTAAGGACTGGTAATGTTGTGAGACAATTAGTAATAGTTGCCTCGTCAAGATATTTAAGTATTTTTGGCCACC
TATTTACAAGGTCAAGAAGGTTATATTATCTTACAGTTCATCTATGTGCACATATCTTTAAATGAGTGATGCTTTTTTT

Fig. 6.336

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TTTTCTACCTTTCTATGGTTTTTATCCCACCTGTTCTCATGCAGTTTTTACAAAAAGGCCACGGCATAACAGCCACTTG
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ACCAACTACAATGACAATAGGTTTATTATCTTTTACTCTTAGAGAATAGTTTATCTTTTATAAAGTATTTATTATTATT
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Fig. 6.332

ATTTAGTATTCAGAGCTTCTTGTTTTTGTTTTGTTATACTTTGTATGTGATAAGGATGAATGATATTTACTCCCTGGCC
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TAGCTGGACTACAGGCGCATGCCACCATGCCCAGCTAATTTTTGTATTTTGAGTAGAGATGGGGTTTCACCATGTTGGC
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TTAGTAAAGCTAATTAACATATCTATATTACCTCACCAATTTATCTTTTTGTGATGAGAATGTCAAAAATCTATTTTAG
AAATTGTAAAACATATAATACGTTATTAATTACTGGGGTCTCC: GCAGTGCAATAGATCACTAAAACTTATTCCTCCA
ATCTAACTGAAATTTTGTACTGTTTGATCGACATCTTCCATTTCTCTCTCCCTCCTCCTCTACAGCCTCTGGTAACCCA
CCTTTCTACTCTCTGTTTCTGAGATCGACTTTTCTAGATTCCCCATAAAGTGAGATCATTTATTAAAGACAATATTTGT
CTTTCTGTGCCTGACTTATCTCACTTAGCATAATGTCCTGTAGTTCCATTCCATGATGTTGTGAATGACAGAATTTCTT
TCTTTTATAAGGCTGTGTAATCTTTCTTTTTTTTTTTTTTTTTTTTTTTTTTGAGGTGGAGTCTCGCTTTGTAGCCCAGG
CTGGAGTGCAGTGGTGTGATCTCGGCTCACTGCAAGCTCCGCCTCCCAGGTTCATGCCATTCTCCTGCCTCAGCCTCCT
GAGTAGCTGGGACTACAGGCGCCCGCTACCACGCCCCACTAATTTTTTGTATTTTTAAGTAGAGATGGGGTTTTACTAT
GTTAGCCAGGATGGTCTCGATCTCCTGACCTTGTGATCKGCCCGCCTCGGCCTCCCAAAGTGCTGGGATTACAGATCTG
AGCCACTGCGCCCGGCCTGTATAATCTATATATATATATATATATATAATATATATATATATATATATATATATATATA
TCACATTTTCTTTATGCACACTTGATGAACACTTAGATTGCTTCCATATCTTGGCTGTTGTAAATAAAGCTGAAATGAA
TATGGGAGTGCAAATATCTTTTTGATGTACCAATTTCAGTTCCTTTGGGTGTATACCTAGAAGTAGGATTGTTGGATTT
TATGGTAGTTCTATTTTTAGATTTTTGAGGAACCTTCATACTATTTTCAATAATGGTGGTTCTAATTTACATTCCCACC
AACAGTGTAGAAGGGCTCCCTTTTCTCCACATCCTTGCCAACACTTGTTATCATTCATCTGTTTGAACATAGCCATTGT
ACATGGTGGAGAGCAACACATACTGGGGTCTGTTGGCGGGTGGGATGGGTGCAGGGAGAGCATCAGGAAGAATGGCTAG
TGGATGCTGGGCTTAGTGCCTAGGTGATGGGATGATACGTGAAGCAAACATGGCACATGTTTACCTATGTAACAAACCT
GCACATCCTGCACATCCTGCACATGTACCCCTGAACTTAAAAGTTGAAGAGAAAAAAAAGGAAAATAGTCATTCTAACA
GGCATGAGGTGATAGCTCATTGTGGTTTTAATTGCATTTCCCTGAGAATTAGAGATATTGTTGAGTTTTCGTTTGTTTT
TGAGACAGGGTCTTGCTCTTTTGTTCAGGCTGGAGTGCAGTGGCGTGATCATGTCTCACTGTAACCTCAAATTCTCGGG
CTCAATGATCCTGCCTCAGCCTCCCAAGTAGCCAGGACTACAGGTGTGTGCCACCATACCTGGCTTTTTATTTTTTCGT
ATAGAGAGGGTCTTGCTATTTTGCCCAGGCTTGTCTAGAATTCCTGGCCTCAAGTGATCTCCTGCCTCAGCCTCCCAAA
GCACTGGGATTMCRGGCATGAGCCACTGTGCAGGGCTGGGGAACATTGTTTTTTTTTTTTTTTTTTTTTTTTGTATCT
CTGTTGGCCATTCATATTTCTTTTTTGAGGAAAGTGTGTCTATTCAGATCCTTTGTCTAATTTTAATCGATTTGTTTTC
TTACTATTTGGTTGTTTGAATTTTTTATATATTTTGAATATTAGCCTCTTATCAGATGTATGGTTTGCAGATATTTTCT
CCTGATCCATGGGTTGTCTTTTCACTCTATTATTTGGTTGCTTGTGCAGGTACTTTTTAGTTTAATGTAGTCCTATTTG
TCTATTTTTGTTTTAGTTGCCTGTGCTTTTGGAGTCCTATCCAAGAAATCATTGCCCAGACCATTGTTGTGGAGATTTT
CCCTTATATTTTCTTCTAGTAGCTTTACAGTTTCAGGTCTTATGTTTAAGCCTTTATATCTCTTTTGAGTTGATTTTTA
TATGGAGTGTGAGATAAGAGTACAGTTTCTTTCTTCTGCAGGTGGACATCCAGTTTTCCCAACACTATTTGAAGAGACT
GTCCTCTCTCCATTGTGTGTTCTTGGACCCTTTGTCAAAAATCAATTGACTGTAAATACTTGGATTTACTTCTGGGCTT
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CAATAGAGGAACTGCATTGAATCTATAGATGGCTTTGGATAGTGTGGACATTTTAACATTACTAATTCTTTCAGTGCAT
GAACATGGGATATCTTTTCATTTATTTCTGTCTTCTTCAGTTGTTTAAATCAATGTCTTGTAGTTTCCAGTGTATAGAT
CTTTCACCTTGTTGGTTAGATTTGCTCCTAAGTTTTTTGTTTTTTTTTTTTTTGTGATGGAGTCTCGCTCTGTTGCC
CAGGCTAGAGTGCAGTGGCGCGATCTCGGCTTACTGCAAGCTCCACCTCCCAGGTTCACGCCGTTCTCCTGCCTCAGCC
TCCCAATTAGCTGGGACTACAGGCGCCTGCCACCACACCTGCCTTTTTTTTTTTTGTATTTTTAAGTAGAGAYGGGGT
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GAATAACAAAATGAGAAAACATTCATCTTGCACCACCTTAAAGCATTTTGCCTACCATTTACTAACACTGGGCATACTT
TTGGACACACTAGTCTAAATCAGCTGCAAGACAATTTAGAGACTAATTTATTGTTTTAAACAATCAACATTTTTCCTT
CTTTACCCTCAGTTGGATTTCTCCACTAGAGGACAGAAAATGGCAAATTCGTTCAGCAGAAATATTCCAATTCATAAAATA
TTCATTCCTTAGGCAGAATGGTCCGTGGATTTAGAGCACCCTACTATAGACAGCCTGCCTGAATTGGATTCATTCTCTG
CCTTTTACCAAACGTAAATTGGGCCAYGTTGCCTTCTGTGCTTCAGTTTTCTCATCTGTTAAATAGTTGCTGTGGTGAT
TTAATGAATTAAAAGATGGGAATTGTCTCGACTGAGTCAGGAATCATACTAGATACTCAGTAGAACTCATTCATTTCCA
TATGTGAAGGAATAACATTAGTTTTTATTTTATTATTTTGGGGGGGTGAGGGTTAGAGAAGGGGTGTCTGCCATCCTGATT

Fig. 6.338

TCTATCCTTTTGATAAAGTGACCTAATAATATTGATACTAACCAAAGAGGACTATTATAATGTTTTATTTACTCTTCTC
CCCCATTAATCACTTTTTTTGGTGGTGTTTTATATGTTTTATTCCTTTATTTTCATCCTTCAAGGAGTTTGCCCTTTGT
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CTCTGCATAAGACTGAAAGTAACCCAGGCACACAATATCTTATTAATATTTAATATTAAGAAAAGTATATTAAGAAGGT
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GAACCAAGACTCTTTGAAAAAAAATAGCTGATTTCTAGGGACTCAGGAAGGAAATATGTAAGATGAGCCTGGAGCATCT
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GGTAAAATAAATATCCATAAACCCATATTGATACAATTTATTAAATAAATACCTAAGTGGGGGAAAATAGACAAATCTC
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CAAGGAAAACCTGAGATACCTACAGCCAAGAAGAGCCTCAGGAGACATGAGGACTAAATATTATGTGGTATCCTGGATG
GGATCCAGAACAGAAAAAAGGAAATTACGTGAAAACTAAGGAAATCTGAATGAAGTGTGGACTTTGTTTAATAATAATG
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TGTAGTTAGTCACATAAATTAGTACAAGAATGCATGATGTTGGTTAATGCAAATGCTGTTTTTTACCTTGTATTGGTAG
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ATTTCTCACTTAATCTCAGTTCTGCTTTTAGATACTCATGTCTCAGAAATTTTATCATAGTATATGCTAGATCAATCTC
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AATTACCCAAAATTACAAAGAAAAGTGCATTTATTATTAAGGTAACTTGTGTTGTCTGTGCCTTTACATCAACTCCAAG
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GGCTCACACCTGTAATCCCAGCACTTTGGGAGGTCGATCACATGAGGCCAGAAGTTTGAGACCAGCCTGGCCAACATGG
CAAAACCTGGCCTGTACTAAAAATACAAAAATTAGCCAGGTGTGGTGGTGCATGTCTGTTATCTCAGCTACTCGGGAGG
CTGAGCAGGAGAATCACTTGAACCCAGGAGGTAGAGGTTGCAGTGAATGGAGATGGCGCCACTGCACTCCAGTCTGGGC
GACTGAGTGAGACTCCATCTCAAAAAAAAAAAAAAAAAAAAGTAGAATTCATGGGAAAAATTATTTGTTGCTATTTAGCC
TTATTTTTAAATATTTCAGTATATTGTCTGTTTGATACATATGAACAATCTGACTACAACTCATTGGGAAACACCAGTA
TTTACCTTACCTTCTAATGTAAGGCATGATTCCAGGTATTTTCTCATACCTCAAACCTTAAATCTCTAATTTAGTCCCA
GAAACAGTATTCTACATGTCAAAACGTTTTTTGTTTTGTTTTGTTTTTGGAGACAAGTTCTCACCTTGTCAACCAGGCT

Fig. 6.339

GGAGTGCAGTGGCGCCATCATGGCTTACTGCAGCCTTGATCTCCCTGGCTCAAGCCATCCTCCCACCTCAGCCTCCTGA
GTGGATGGAGCCACAGGCATGTGCCACCAGACCGGCTTTTCTGTTTTTTAATTTAATTTAACTTTTTTTTTTTTTTTTT
GTAGAGACAGGGTCTCACTATGTTACCCAGGCTGGTCTCAAACTCCTGGGCTCAAGTGATCCTCCTGCCTTGGCCTCCC
AAAGTGCATGAGCCAGTGCACCCAGCCTCAGAATATTTTTAAAGATGAAAACCTAATGCTCAGTTAAACTTTTATAAGA
TCTATGTGCTTCTAAATAAGTGTAAAGATGACTATTTTCTGGATGTTGTATTGGTGGGGAAGATGGAGAGATAGAAGAA
CAAGTGTTTATTCTCTGGTTGTATTTTCTCTCTCTCTCTCTTTTTTTTTTTTTCCTTGAGTTAGAGTCTTGCACTGTTGC
CTGGACTGGAGTGCAATGGCGTGATCTAGGCTCACTGCAACCTTCTCCTCCCAGGTTCAAGCAATTCTCCTGCCTCAGC
TTCCCAAGCAGCTGGGATTACAGGCGCCTGCCACCACGCCTGGCTAATTTTTTGTATTTTTAGTAGAGACAGGGTTTCA
CTATGTTGGCCAGGCTGGTCTCGAACTCCTGACCTCGTAATCTGCCCACCTTGGCCCGCCAAAGTGTTGGGATTACAGG
CGTAAGCCACTGTGCCCGACCTCTGGTTGTATTTTCAAAATATATCCAAATAATATTTTAAAAATGTATTTAGTGAGCA
CTAGATTCCAAAATAGCAAGAGCAATTTCAGCAAAGTATAATTCCTAGAGAGGAATCCTACAGTACCTCGTTTTGCCTT
TTTTGTTCTCTAGGTCATGTCAGGATGGGAGCATAACTGACCTGGCCCTAGTTCCTAGGAGGAGCATGTGCCCTCTAAC
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GGATTTGGAGGTGGGATGGAGGGATAAAACCTTAGTGGGAAGCAGACAGCAGATAGAGAAGATATAGGATAAACTCACG
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TATTAGGCAGTGTGAGGCAGAGAAAGAAGAAGTGGGAGAAGGAGGGGAGTCAGACTTTCCTTGGCTGGAGGCAAACTG
ACCAATGCCCGTTGCTGCCCCTTTCCACCGCAGGGTTCATTGGTGAAACCTGGTTGGACTGAACCTTTAGGAGAACCT
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GGAGGGAGGGGTAGAGCTGACTGGAGAGCGATTTATGGATGGTATATTTCATGAGGCTTGAGTCTTTTTTAGCACTAAT
AGTTGGCGTCAGCTTCACCTGTTGCCTCCATGAGTCATGCCCTGGTAGCAGAGGATGGTGGGCCAACTGCCAAGCCATC
CCCTGAAGGACCGGCTGCCTGGGAGTGAGGAAAGGTGGTTGATGGGCTCTGAGACAGCAAGGTACAAACTGAAATGGGG
TGAAAAGAGACTGTCAAAATAAGTATGGGCTGATTTGTTCTAATATATCATAGGTTATTATTAGATGCTGGAAGAGTAA
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TCCTCTTCTGGATATAAATCTGCCCTACAGTAGAGTTGCAACCTTGGATAAATAAATAAATTTGGCCTCCCTGAGCCTC
TGTTTTCCCTGTAAATCAGGACTAACATAATCTACCTCAAAGGATTGTGTTGAAAATTAAGTGAAAACTTACATAAATT
ATGTAGTGTGGTGGGGACATATGATAGGTACTGTGGCCTATCCTTAGGAGGGATGAGGAATGGAACTATTTTTAAACAT
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AGGTCTTCACTTCACTGTGTCTAAACATTGGTCTGTTGCTTGTGCTTTGCTTTTCTCTCACCCTGGATGACCTTCAAGG
CTCTTCCAAATGATTTCTCTGATTCCCTGAAATACTGAAATTGTGATTAACAGAAGTTTCAAGTGTTAATCAATATCAA
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TGAGTCTGACGCTAGCCACAGCCGGTAAGAGAAGGAATTCCCTAGTAGTGTAGTTGTTACAAGGGAATGAGTCATTTTG
CAGGTAATTGTTCTTCTTCCATTTTCCACTTGCTGTCAGCCACTTTACAGAGGAGTCGGACTTTGTTCTGCTTAAGAGA
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ATTCTCCTGCCTCAGCCTCCTGAGTAGTGGGTATAACAGGTGCATGACCACGCCCAGCTGTTTTTTGTGTTTTTAGTAG
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AATTATTGTATGACATGTATCTGCTCCAGAGGTTCTTATAGGTGAAATGAGATTTTTATGTATCTGTATTCAAACACAT
AGATAGATAGATAGATAGATAGATAGATAGATAGCGAAACATATAGACATACCTCTGAAAATACAACCAGCCATA
TAATGTAAGCCTGTTGTGTGTTAATGAATGATAGGGTAATAATACATTATGTTCCTGTCCTGTAGAGGAGTTCGAGTAT
TCAATACAGAGGACCAGAATGATTGGTCCCAGAAAAAGTGCTCCTTAAGCTGGATCTTGAATACTGAGGATTTATATAG
GCATCTAATAGGGAAGAGGTGGTATTCCAGGGGCTGGATAAGGCAGGGTAGGATGGGGTATTCATGTGGATTCGCTTTA
TTGTATGTTTGCTGGAGAGGAGGGTAGTGGGCATTCTGGAGGAGTGAGCCTCTCTGGAACAGAGGATGTGATTATAGGA
CCAGTGGCAGGTGTGAGAAATGTAGATTGAGGCCATACTGCAGACTGTCTTAAATGCCAGGTCAAGCCATTTGCACATT
GTCCTATAATTCATGAAGAACCCTTACATTTGACTATAAGGAAAATATTATGAAGAAACTAAAATAAGGTGACTTTAGA
AGGTTATATCTGGTGGTAGTGTGCAGGATAGATTATATCAAGGGGAAATTGTACTGAAAGCAAGAAGAAACCGTTAGGR
AATTAGGTGTAAGAGTGAATAGGGCCTGAAATAGGATAGTGATGAAAAGAGAGGAATGAGTAGGAGAAAGATTGCACAG
AGAAGGGTGACAGCATACATAAGACTTGGCAGGACCACAGCCAGAAGAAAGCATGAAATTCTAAACCTGAATAATGGAC
AGACATATAATTTAGGTAAGGCTATGTATCTAAGCGGAAATGCCCTATAAGCTTTCAGTTGTCAGGACTGGATTTCTGG
TTAAAAAAAAAAAAAAAAAGAGGGGGGCAGGATTTTACTTGGGTGTTGTCTGCAAAGTTGATAGCTAAGACCTTAAGGCT

Fig. 6.340

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CATCTGTGAAACGGGAGTGACAGTGCTCATTAAAGGTTATTATTAGAAATAAATGAAGTAATTCCAGGAGAGTACATAG
GACTCTCCTTGACACATAGTAGACCCTCAATAAACATTAGGTTTTATTATTGTTTATTATGGGTTTTAGGGAACAACAC
TTTTGTGAATTATTAGAACCTTATTTGGGAAGCTATAAAAGCCATCTGCGTGATGTGTTTCTTAGTACACCAAATGGA
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GTTTGTAAGGCTTTTCGTGTTTTTTACATATTTGGTTAAGGATCTTAATCCTTGAGCTGTATTTTGTTCTCATTTGTTT
GTGTCCAGAGTCAAATTAAGTTTAAGGAATTTATAAAGAGCTTCTTTGCAAGAATGATATTAACTCAATGTACTAAAGC
TACATTCACCCAAAGTGTAATAGACTTACACAACATGGAAGAAATCCAGGAGAGGCAGACAGTTATTCATGAAGAGTGG
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AGCATTCTCAGCCAGTATCCGTGATTCTGATAGTTTACAAACTGTTGATAGCATAAGCTGTTAGGAAGATATTGTAGAA
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CTCCTGACACATGACACTGCTGGATCACCAACCTTAACTCTGATTCTCTCCATACAACCCAGTATCCTTAAGTAAAATG
CTGCATATTTCATTGCTGGTAAGAGAGGCAGGGCCAAGATTCGAACTCACATGAAGCCAACTAGAGAAGCCTATGCTCT
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TTAATTGCACTTATTATTGCCTTCAGTAAAGTCTTCATTTCTAAAATGTCTTAGTAAAACAGTAATTTCATAAAAATAT
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GTTTGAAAGTTACTACTTATTCAAAACTCAACTGTCTGGCAGCTTTTCTTATCCTGAACTCTGTGAAGAACTTCGGGGA
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CTTCTTTCTCCTTACATACCCTCCTTTAAATCCGTACCTCCCCCGGTCTTCTCCATCTTGCAAATGGCACCAATGTCCA
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TGCCCCAAATCTGGTTGGCTTATTCTAGGCATTCGAGTCTCTTCCCAAATACCACCTTCTCAAGTGCAGCTTTTCCTTA
CTACTTAATTTTATAGTCCCCCATCAATGTACGATGTTACCCTCTTTTCCTTCATTGAGAGTGTTTGTAATTTTTTTTT
GTCTGCTTRTTTATTGTCTTTTTGCCCCATAAGTCCAAGGTTTACATGAACAGGGAACTTGTCTGTTTTGTTTATTACT
GTATCCCCTATGCTGGCACATACTATGTAATAAGTGTTTGTTGAGTGCATGAGTGAATAACCATTCTAAAAAACTCTGA
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CATTGCTTTTGGTGTTTCAGACATGAAGTCCTTGCCCATGCCTATGTCCTGAATGGTATTGCTTAGGTTTTCTTCTAGG
GTTTTTATGGTTTTAGGACTAACATGTAAGTCTTTAATCCGTCTTGAATTAATTTTTGTATAAAGTGTAAGGAAGGGAT
CCAGTTTCAGCTTTCTACATATGGCTAGCCAGTTTTCCCAGCACCATTTATTAAATAGGGAATCCTTTCCCCATTTCTT
GTTTTTGTCAGGTTTGTCAAAGATCAGATGGTTGTAGACATGTGGTATTATTTCTGAGGGCTCTATTCTGTTCCATTGG
CCTATATCTCTGTTCTTGTACCAGTACCATGCTATTTTGGTTACTGTAGCCTTGTAGTATAGTTTGAAGTCAGGTAGCA
TGATGCCTCCAGCTTTGTTCTTTTTGCTTAGGATTGTCTTGGCAATGCGGGCTCTTTTTTGGTTCCATATGAACTTTAA
AGTAGTTTTTCCAATTCTGTGAAGAAAGTCATTGGTAGATTGATGGGGATGGCATTGAATCTATAAATTACCTTGGGCA
GTATGGCCATTTTCATGATATTGATTCTTCCTATCTATAAGCTTTGTGTCCTCTTTTATTTGTTGAGCAGTGGTTTGT
AATTCTCCCTGAAAAGGTCCTTCACATCCCTTGTAAGTTGGATTCCTTGGTATTTATTCTCTTTGAAGTAATTGTGAA
TGGGTGTTCACTCATGATTTGGCTGTTTGTCTGTTATTGGTGTATAGGAATGCTTGTGATTTTTGCACATTGGTTTTGT
ATCTTGAGACTTTGCTGAAGTTGTTTATCAGCTTAAGGAGATTTTGGGCTGAGATGATGGGGTTTTCTAAATATATAAT
CATGTCATCTGCAAACAGGGACAATTTGACTTCCTCTTTTCCTAATTGAATACCCTTTGTTTCTTTCTCCTGCCTGATT
GCCCTGGCCAGAACTTTCAACACTATGTTGAATAGGAGCGGTGAGAGAAGGCATCCCTGTCTTGTGCCAGTGTTCAAAG
GGAATGCTTCCAGTTTTTGCCCATTCAGTATGATATTGRCTGTGGGTTTGTCATAAATAGCTCTTACTATTTTGAGATA

Fig. 6.341

CATCCCATCAATACCGAATTTATTGAGAGTTTTTAGCATGAAGTCCTGTTGAATTTTGTCAAAGGCCTTTTCTGCATCT
ATTGAGATAATCATGTGGTTTTTGTCTTTGGTTCTTGTTATATGATGGATTACGTTTATTGATTTGCATATGTTGAAGC
AGCCTTGCATCCCAGGGATGAAGCCCACTTGATTAGGGTGGACAAGCTTTTTGATGTGCTGCTGGATTTGGTTTGCCAG
TATTTTATTAAGGATTTTTGCATCGATGTTCATCATGGATGTTGGTCTAAAATTCTCTTTTTTTTGTTGTGTCTCCGCC
AGGCATTGGTATCAGGATGATGCTGGCCTCATCAAATGAGTTAGGGAGGATTCCCTCTTTTTCTATTGATTGGAATAGT
TTCAGAAGGAATGTTACCAACTCCTCCTTTGTACCTCTGGTAGAATTCAGCTGTGAATCTGTCTGGTCCTGGACTTTTTT
TGGTTGGTAGGCTCTTAATTATTGCCTTAATTTCAGAACCTGTTATTGGTCTATTCAGGGATTCAACTTCTTCCTGATT
TAGTCTTGGGAGGGTGCATGTGTCCAGGAATTTATCCATTTCTTCTAGATTTTCTAGTTTATTTGTGTAGAGGTGTTAT
TCTCTGATGGTAGTTTGTATCTCTGGGGGATTGGTGGTGGTATCCCCTTTATCATTTTTTATTGCATCTATTTGATTCT
TCTCTCATTTCTTCTTTATTAGTCTTGCTAGTGGTCTATCAATTTTGTTGATCTTTTCAAAAAACCAGCTCCTGGACTC
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AGCTTTTGAATGTATTTGCTCTTGTCTTTTCTAGTTCGTTTAATTGTGATGTTAGGGTGTCAATTTTAGATCTTTCCTGC
TTTCTCTTGTGGGCACTTAGTGCTATAAATTTCCCTCTACACACTGCTTTAGAATGTGTCACAGAGATTCTAGTATGTT
GTGTCTTTGTTCTCAKYGGTTTCAAAGAACATCTTTATTTCTGCCTTCATCGCATTATGTACCCAGTAGTSATTCAGGA
GCAGGTTGTTCAGTTTCCATGTAGTTGAGTGGTTTTGAATGAGTTTCTTAATCCCAACTTCTACTTTGCACTGTGGTCT
GAGAGAAAATTTGTTATAATTTCTGTTCTATTACATTTGCTGAGGAGTGCTTTACTTCCAACTATGTGGTCAGTTTTGG
AATAACTGTGATGTGGTGCTGAGAAGAATGTATGTTCTGTTGATTTGGGGTGGAGAGTTCTGTAGATGTCTATTAGGTC
CGCTTGTTGCAGAGCTGAGTTCAATTCCTGGATATCCTTGTTAATTTTCTGTCTCGTTGATCTGTCTAATGTTGACAGT
GGGGTGTTAAAGTCTCCCATTATTATTGTGTAGAAGTCTAAGTCTCTTAGTAGGTCTCTAAGGACTTGCTTTATGAATC
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TCTCCTGAATACAGCGCACTGATGGGTCTTGACTCTTTATCCAATTTGCTAGTCTGTGTTTTTTAATTGGAACATTTAG
CCCATTTACATATAAGGTTAATATTGTTATGTGGGAATTTGATCCTGTCTTTATGATGTTAGCTGGTTATTTTGCCCAT
TAGTTGATGCAGTTTCTTCCTAGCCTTGATGGTCTTTACAATTTGGCATGTTTTTGCAGTAGCTGGTACTGGTTGTTCC
TTTCCATGTTTAGTGCTTCCTTCAGGAGCTCTTGTAAGGCAGGCCTGGTGGTGACAAAATCTCAGCATTTGCTTGTCTG
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TTTCAGGTACACCAATCAGATGTAGATTGGTCTTTTCACATAGTCCCATATTTCTTGGAGGCTTTGTTCATTTATTTT
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TCTTCCAGTTGATGGAATTGGCTACTGAAACTTGTGAATGCATCATGTAGTTCTCATGCCATGGTTTTCAGCTCCATCA
GGTCATTTAAGGTCTTCTCTATGCTGGTTATTCTAGTTAGCCATTTGTCTAATCTTTTTTCAAGGTTTTTAGCTTCTTT
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CTCAGATCTCAAACTTCATGCTGGGAGATCCACTACTCTTTTCAAAGCTCAGTTGGAAATGCAGAAATCACCCGTCTTC
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GCTCAGGCAATCTGCCTGCCTCGGCCTCCTAAAGTGCTAGGATTACAGGCATGAGCCACAGCGCCCTGCCAACCTAAAG
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ATGGTTTTGCCTTGCCAGAATGAATAATACATTAAAGATATAGCCTTTTGTGAATGGCTTCTTTCACTTACAATACTTT
TGAGTTTTGAGTTGAATTATAAGTTTCACTTATAATACATTTTGTGTTATTGCATCTATTGGTAATTTGTTTCATTTTA

Fig. 6.342

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TTGCTTTTTTAGTATTTCATTTTTTTCCAGTATGTCATTTTATGGACACAATTTGTTTACCCATTCACCAGTTGACTGAT
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TCATTTCTCATGGGTAGATACTTAGAAGTAGGATTGCTGGGTCATATGGTCACTCTACTTTTTAACTTTATAAGAAACT
GTCAAACCTTTTTCCAAAGTTTCTATACCATTTTGCATTCTCACTAGCAATGTATGAGAATTTAATTTGCTCTGCATCC
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GATGCTCAATTGTTCAAGCATTCTTTGTTGAAAAGAATATCATTTCTCTTTTATAGCTCAAATTTTATTACTTAAAATT
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GGCAGCTGTGTGTTGTTCACATAGCGCTCCCAAACATAAGCGTGTCTCACTATATGGCAGGGCTGTCTGCCTGTGGGCA
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TGCAGATTTTAGAGGGAAGGGAAACTGTATGTGTGTTTCTGCATAATGTTTTAAGACAAGGAGTATTATCTACTATATG
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AAATATTGATGTTTTGCATCTGCATCAGCCAACAATCTTTTTGTGGCAATAGTACACTACCTTGAGAAATACAGGGACA
TAATAAGATACTTTTCTGGCCTCAAAGATCCTCCACCCTGGATGAGGAGTCAAAGTTGAAGACAGTTGAAAAAAATAGA
ACAGTGATTCTTAACACTGGGGAACATTCAGGAAATTTCAGGGGACAGTGACTGGGGAGTACTCCTGGTGCTTGGTGGG
CAGAGGCCAGGGAGGCTAGATGTCCTGGGATACATGGGACAGCCTTTCACCAGGGAGGTTTGCTGTGTGTCCCACACAA
CTTCCAAAAAGTCCCAGCAGCTATTCACATAGGTGAAAAAACAAAATGCATTTATAATTCTCTGAGCCTAGAACCCAAC
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Fig. 6. 343

TACTTTTTACGTATAAGTATTAAATATTTTTACATGGTTTTAATCCACACCAAATTTTCTAGTAATGCAGCAATAGTGT
ATATTGTGAAGATGGCACTTTGTTACATTTGAAACCTTTCAAAGAATCATTCATCATATCAGGAAAAAGGAAAATTTTT
CTAACATAACACCCCTGTATTCATTTTCATGCTACTGTTACTTTTGTGGTAATTCTCCCCATAAGAGAAATCATAGCAA
ACCCTTAGATAGCACTTACTGTTCTAAGAGCCTTATGTATATGCACTAATTTAATCCTCACAACAACCCTGTAAGGTAG
AAACTATTATTCTCATTTTCCATGTGAGTAAACTGAAGTATGGGAAATTTAAATAGCTTCCCCAAGGTCACACAGCTAA
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TACTCATTTCTCCTTTATTTCACTGATAAGCCATTATACTGATTTTTTTTTTAAATCTATGTAGGGAGTATTTAGCCTA
TATGTATTCAGTACAGTATTATGAAGGAAGTGTATTTAGGAATTTCATTTCAGGATGAATACTTGTTATAAAAAGTAAA
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CAGTGAGCCTGACTGCAGAGTGTAATTTGCATTTAAATGACAAGTGGTATGACACTTCCACCTGTTCTTCAGGGAGCCT
CACCAAGTATGTTTTGCCTCGAGGCCTTTGCTTTTGCCATTCCTCACTTGGTTTTCCCTTCTCTACCCTACCTCTCACC
CCGCAATCCACATGGCTTGTTTCCTTTCTTCAGATCTCCTGGTATACGTCATCTCATCAGAGGGGCTTTCCATGATCAC
CTTATAGCAACCTCTATCCGTAATCCTTATCCCTTCTTTCTCTCCATTGTGCTTATCACCCCATGACGTATTATGTATT
TATTAACTTAGTTTTGGAATATAAGCTGCAGGTGAGAAAGCTTTGTTTCTCTTTCTCAGGCAGTGCCTGACATAGAGTA
GGAGCTTAATAAATATTGAATGAAGGGGCAGATGTAGAATCGTTAGAATTTGGAAAAAGGCTGGAATATTTGGTCCTAA
GATGTACTTTGAATGATTTAAATCTGACTGAGAAAACCAGTTTGAGGTTGGCAGAGTAGAAGAAGGAGGGTATTTTAGG
CCGAGAATTTGGTAAGACAGAGATGGGATTGTCAGCTGTATGGTGAAGGTGGAGGARTACTAGTCTTTCAGGAACAGAG
CAAATGTAATAGAGCAGCTGAAAATAGGCTTGGCGTTGGGGATTAAGGTCAGATATGGCAGTCTGGAATTTACAGAGGG
ATGTGATGAAGCAGGCATTACTAGCTTTGAAGTAACAGGCTGCGGCTGGATTTGAGGAAAATTGATCTGGCAGTAGAGG
TCAATATGAGACTAGAAGGGGAAAAGCCAGGGGCAAGAAGATGAGCTAAGAAGCAATTACAGCATTTTGGCCCTGAGAC
ATGGAGTGCTGTGATGCAATTGAAAGGAAATATAACATATATATTTTAAAAACAAATGACAAACCAGAGAATTTTTTTA
AATATTAGTATAATAGAATGATTATCATAATGATCATTGTCACTGTCATCTTTACTGTCTGATACCTACTTATCTTCAA
CTTTGTACCATTTGTTGTTTTAAGTGCTTTGCATATATGGTCTCATTTAATCCTCATATGTATCTGTTCTGGTTATTTA
TTGCTATGTAAGAAATCACTTCAAACCTAATACTTTGAAATATGAATGTATTTTTATTAGTTTTCATGGCTTTGTGGGT
TAACTGGTCTTGGCTAGACATTCTTGCTTGGCATTCCTCATATGGTTGCAGCCCAGTGGGTAGCTATCTGAAAGCTGAA
CTGGCTGGATGGCTAGGATGGCTTCTTGCCCCTCAGGTCTGGCTCTGGGCTGGGGTGGCTGGCATTGCTGAAGACTGGC
CAGGCATGTGTGCTCTTTCTTGTGCAAGTGCTCTCTTGCGCTTTCTCTCTTCACTGTCTGTGAAGTTAACATGGACTTC
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CATTTTTGGAGCAGTTTTTATTTGGAAGGAAGGGAAGCCAGAGGACATTAAAATTCATAGAAAATGCCCTCCAAAAGGA
TGGGCAATTCTTAAGAATGAACTACGAATTTTGAGGAGTTTTATGGTCATTATTCATCACTGCAAGAGGGAAGCCCCGT
TCATCTTTTCTCAGGGTGGCCTCAGAGCTGCAGGGATACCATTATTCAATGGCGTTTGTCGGGTGAGGAGAAAGCCTCC
CCAGAGGCTGGCCCTTGCCAACCAATCCCAAAGCAGCCTGCACCGAGGCCACACCCCCTCGCCTTATAGGCTAAGAGCT
GGAATGCAATTGGTGCAGAGTTGGGGGTTTTATGGGAGGGGCTTCTTGTCACTCTTCCCGGCTTCCCTGCAGTTCCTTAT

Fig. 6.344_

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TTGGTAGCTTTTGACAGGACTAGCCTTTCTTGCAACTAAGCATCTTGACATACATTATTCATTAAGCCCTGGAGCTCGG
GAGAGAAAGATGCAGACCCTTAGATCTTTAGATATTCCTTTATCACGTGGATTTTCTTTATTCAGAATAGTTGCTGAAT
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TGAAGCTAAATACATTTTAGAGCTKWCTATATATATACATATATATATATATATACATATAATCAATCAAAAATGCCTG
AAGCAAACTATTTACTGTCAGTGTCTTGGGGCTACATAAAGGTAAGACTTACTTAGCTTTTGAAAACTCCTTACTCTGA
TTACAAAATTATTTAGAAAGCTTAGACTCTACTGTAATTTGTTCAAACTATCAGTTATGTATTCTTTCCTTACACATGA
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GCTKTTTTAGATACGCAGTAGTTCTTCTGTTTGTCATGTGGTATATGTTTGAATTGTGCTTGAAAAATATATGGTAATT
AAATACATTCATTGCAAATAAATTATTGGGCCAAATTACTTATACTGATTTATGGATTCCAGTTAGTATGTTGCACATA
AAGTTTATAAAATTAATTTGTGGCTGTTTCTAAAAATCTATACTACTTTAACCTGACGAGGAATACGTTTTTTTCACCT
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GTTTGCTTTGTTTCTTTCTTTTTGAAGGACAAGTTCCCCTTTGTTTTAGAGATTTACTTGAATTCTAAAAAAATTAGAA
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ATTTTAGTTTGCTTCCTGAACAATCTTAAATGCCTAATGTAAATTGAAGAATTGCAGTTCTGAAAAGCAAAATACAGTA
TTGAGATTCAACTGCATTTTTACTTTCCTTTATGCCTTAACTGCTGTACACAGACATTCTGATGTATAATGAGAACAAA
GGATTCAAAAGCATTCACTTAGAAATCCTCCCCCTGTTTTTTTAGTTGCAACCCTAAATCTGTGTATTGTTTTCAGACTA
CTTAGGCCAAAAACAATTAGAAATTCATCAATGGAAAATATTTAGAGGTCACTTAAAAAAATAAACTAAATGCTTAAAT
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AAAATACCCTTGTGTTATAACTTAATGTACGCTTCAGAATTATCTTTAGGAAATTCCTTAGACCGTCTTCCTAGAGTAG
AGAAGTAATTGCTTCAAATATTGTCTTTTATAATTATGTTAAAATGAAATGTTGACTTCCTTGGAGTCCCTTATAAGCC
TTGGTAGGGAGGTGGGCATGTGATGGAGGATTTCTCCAATCCATGTTTTTGTGTTTTAAACAAAGGCTGGAAAGTACTC
TGGGAATAATGTATATGACCAGAAAGATGAACATGCAGGATGTCACTTATCTAGTCTGTACAATATTTAGATTCCTTTC
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TGAACTCATCCCTTTTTATGGCTACGTAGTATTCCATGGTATATATGTGCCACATTTTCTTAATCCAGTCTATCACTGA
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GTAACTATTTTTATGGTTATAACTACAGTCAAATCCTTCATCGTCTGACCTGCAATGATAAAAATCAGCTATTTACTT
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TGTGCCAGGCTAAATAGAAGACACTTCTATATTAGCYCATTAAAATTATATGATAGCCCATAATTTACTCAAGAAAATAT
AACTTTGTAAAGAGGGACAGAAAAAATTTTGAACTCTATTATAAATGTCTACAAATATTCTTAGAAGGCCCAAAGTTTA
TTTTTTTCAGTAGGTTATAAGATATAATGCTGAGTGAACACAAGCAGTAACCTATGTTCTGTATACCACCTGATGCCAG
TTTTAAAAATATGTATTCACATACAAGGGTAGAAAAAAGGCATAAAAGGAAATTTAACAAATTATCTGTGGTTATCTTC
CAGCAGTGGGGATTTACATTATTTTCTTTTTTTTGGTATAGCTTTCTACATTTTCTATAGCACATGTACATTTTATAAT
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Fig. 6.345

TAGAAAGGAGTTATAAAAACATATGAACAAAGAAGTGAATTTAGTGTTCCTGAGTTTGAGACTAAGCTTTCTAGACCAG
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AAGAATTATATTTCAAACATCACTAAAATATACATTCAACACTTACCTTCATACTTAAATGAGAATTTATGGTGAAATT
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TCAGAGTCAAAGGACTCTGAAATGCTTAATGCTTAAGAAACAGCTGATGGGGGTGGGAGAAGTGTAAATCTTTGGGAG
ATTTGGATACTGGTAATGCCATGTGGTGTTGGTAGGTTAATTCCAGTGTTTCTTTCAATGAAATAGGCACTATGGCATC
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GCCCTTGATCAGTATATATATTCATTTCTGCTAATGATAAATTAATTGAAGAAAGCATTCCATGTTTTATTGTATGGCA
ACACTTTTATGTGCTATAAATTAGTTTCTCTCTTCTCTAAATAACTAATAATACATCTTAATTTTTAATTCCCTTTTCT
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GCCACCTAAGGTCTGTAATTTTTAAATGATTGTGTTCATTGAGTCATATTTTACTCCACTTTCTAATATTTTAGTATCT
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<div align="center">Fig. 6. 346</div>

TACTAACCTTAGAATATAGTTTTTTCAGTCACTTTCTTGTGTCCTGCTTCTTTCTTCATATCCATTTCTCTCTCTTATT
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GGAGGAAACTGAAGAAGATAATTTTATCTCCTTAGGATGTCTGTTTAAAGTCATTTTGCTGGTGACCTGCCTGGGATGG
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CAGCGACATAGGGAAGTTCACAGTTGCCAGAGTAGCTTTGGATTGCTAAAGTTTTTTTGACGATGAGGTGTGATGAGGC
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ATTATCAAAATTACAGCCTTAGTTAGTTTGGTTTGGCTAGGGATCATGTAAGAGAATTATCTTCCCAGCATGCAGTAAA
GGAATCCTTCTAATAACTTGTAAACTTGTGATATGTAGCTTCGTGAAATATTTTATCAAAATTTGTGCTTATTTTTAGT
TTGCAGTAAACCTTTTTTAAAATTTGACTTTTTATATTTTATGGATGGCTTGAGCATCCATGTGTCAAGCCAGCACATT

Fig. 6.347

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CTCAGCTCTTGCCTCAGAGCTGGAGCTGCCATCCTGTCCAAAGCCTGCAGCTGAATCCATATTTCTCATAATAAAGAAT
TCTAAAGACCTCTGATTATCAAATTTATAAACCCATAGTTGGTTACTTGTCTTACTTTAAGGAAGCTACGGAAGCACTG
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GTTGAAAGCATGCTGGAGGTTCTGCAAGCAGAGAGAACAATTCTACCTGGTAGAGTTGGTTAAGCTATAATAAATGATT
TGAGTGTGGGTACTGTGAACAGGATTTAGAGAAATTGAAATTTCAGGCAGGAAGCTGCGTAAATATTTTTAAAGGATGA
TGCATATAAATAAATCCATAGGCCAGGTGCAGTGGCTCACTCCTATAATCCCAGCACTTTGGGAGGTCAAGATGGGTGA
ATCACTGACGTCAGGAGTTCGGAGACCAGCCTGGCCAACAAGGTGAAACCCTGTCTTTACAAAAAAAATACAAAAAAA
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ACAGTGAGTGTAAGTCTTACTTACAGGATACATATCTATAGTGTCTGCCCCAGTCTTAACTGTTTCAGCTCCAGGTCTT
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GGGAGGCAATGTATTCCTTCTTTGTGTGTGTTACTCTGGCCCATTTACTAAGTGATCTGAAAAGCTGCTAGTTTAGATG
GGGCACAGAACAAGAGAGTGTCACAAAAACTGCTAGTTTTGAGTGGGGCTCTACAACAGGTCCAGCCTGCTGTGCAAGC
TGCTCTGCACATGGGCCACATGATCCAGCAGATTTAATGGTGCTTGAAATGTCAGTGGCAGATAGGAATGTTGTTTGGA
GCCTTTGACAGAGCCCTCTGTGTGAATCACAGCACAGACTCTGGAGCAAGACCCTGCCATCATCCAAAGAAAGATAACT
AACTACTCTCCTTTTGAAAGAAAGATATAGGCCTGCTGCTGGGCCTTAGGAGATACTGACTGACTATGGGCCACCAAGT
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GAAAAGGCACTTTCTCTGGAAACAGATGTGCCTTCCCTGCAGTTTTTCTGCCAAAACTATGGTATTCCATACAGCATTG
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ATGCCCTGTAATTAAGGTCAGCACAAAATAACCCAATCCAGGCAGGACCACTAATAACCCATACCCTTCACGAATGAAG
GTTTCGGTCACCCCACCAGGTAAAGAATCATGACCAGCTGAGGTGCTTGCTGAGGGAAAAGGGAATACAGAATGAGTAG
TAAAAGAAGGTAATTACAAATACCAGCTATGACCATATGACCAGTTATAGAAATAAGGACTATAATTGTCATGAGTATT
TTCTTATGAATGCATTTATATGTATATATACATATATTAAGCATATATCTTCATTTTCTTTTTCTTATTCCCTTATATA
ACATAAGAGGTATTAACTTATCTTCATTTTCTTTTTCTTATTCCCTTATATAACATAAGAGGTATTAACTTTATATTAG
TATTTAAGTATTTATTTTATATCATAGTATTTAAGTTATAGGCTATCAGGATAAGAGTAAACATTACTCAAAAACTTTA
CTTTCACTTCTGGGGAATGTGTTAGTGTGCTTTTAGTTGTATGCAGGATAGTTGTAGCCTGTTTGGTAGAATTATGGCC
TTATGGAGATTAAATATGGTTAAAGGAGATGCTTATGGGTACCAGGGTGACAAGGGGCAGAATTTGTAATGGTTAATTT
TATGTGTCAGCTTGACTAGGCTAAGGGATGCCCAGATAGCTGGTAAAACACTATTTTGGGGTGCGTCTGTGAGAGTGTT
CCTGGAAGAGATTAGATTAGCATTTGAATTGGTAGACTGATTAAAGAAGATTGCCCTTACCATTGTTGGCAGGGATCAG
TCAATCCATTGAGGACCTCAAAGAGAAGAACAAAAACATTGGGGGAAGGGGCAAATTTTGCTTTCTTCGGAGCCTGGAC
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Fig. 6. 348

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ATACATATTCTCCTGCCTCTAACATCAGGGTTCCTGGTTGATTCTCTGGCCTTTGGACTTGCTTTCCTGGTTCACCTTT
GCAGACATGGGACTTCTTGGACTCCATAATTGCCTGAGCCAATTCCTATAATAAATAAATGTGTCTTTTTCTCACATAC
ACACACACACTCATACATATGTCTATTCTATTTCTCTGGAGAACCATGATTAATACAGAGAGAGAAAAGATTAGCACAG
ATGAAGTACATGTTCTCAGTTATGTGGTAGTACCTGCCCACCCTTCCCCCCATTTCATTAGCACTCAGAAGAGAGGGAC
ACAAAGTGGTCTTCCTGCCTTCAGTAGTAGCATATGTTGGGCATAATTTAATTTATTCTTGATGATCCAGGGTAGTTG
TAACAAATGAGCACAATTGATCTATATATAATAAAATGATGGCTTTGAGTTTGTAAAGGTATGCATGGCCTCTCAATAA
AAAATAAATACTTACAAAGTTGTCTTTATAAATGTGTGCCAGGCACTGAGTGGACTGTGTTGATTTCCTGGTTCATGTT
TAGTGTTCACTGCTCAGGTTTTCACCTCTATAAGGTACTTGTAATCATAGTCAGTATAAGGTGAGGGCTCTAGAAACTG
CCTGGTTTTAAGTGCTGGCCCCAGCACTTACTGGCTAGGTGATGTTGGCAAAGTTACCTACCTTCTCTGAGGTACATTT
TCTTCATTTGTACAGGATTATATGAGTATGTCTATGTAAATATTTATGTGTATAAAAGATGTCCCCAAGGGACATTTTC
TATCCCCCAGCCTATCCCAAGGACACAAAGACTTACTTCCCACCTATATGGCTCCATGCCATCCATGAATGGAACATAG
CCTTAAAATGTCGATAACAACAGTATCTTCCTAAAAGAGTTATGAGGATTAAATGAGATGATTCACATAAACCATTTAA
CACAATGGCACCTAAATCCTCTAAATGTTGTGCCTTGCTGTATTCCTGTTTGTACTTTGTAAGTTTGAAATAATTGAAG
TGCCTAGGGGAAGCCAGATAACCAAACATATTAAATTAATAACCTTTAATATTTTGTAGGGGCTTAAGTCTCTAAGTAG
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TGGGTGGTACAAAGTAGTATAGAAGTTTACAGGAGAAAGTGGGCAGTTTCATCTGGTAATTAAGAAAGACTTGATGGAA
GAGGTAACTTATATTGGGCTTGAAGATGAAAGGGTTTCAACAAAAATGTTATATAAGCTAACCTCTTTGCCCCTCTGTG
CTGTGCACTGTACCATCCTGACAGCAACTTTCTGATCATTCCTGAACCTTCAAGGACTCTACTTACAAGTAATGGATTA
GTGTCTTTGATGAAAATCTGCTGAGGAGCTGCAGACTCCTACCTCCCAATTTAAATGTGACCATATGCCTTCAGTCCTA
AAGAAGAGTAGAAAGTTAAATAACTTCCTTGAGATTCAGTTTCTTAAATGCTAACATTTGTTCATTTAAAAATCAACAG
TCACCACCACTTTCCTGTAACAGCATCTGAGATGGAAGAGGCTATGGAGGCCTGCCAGTCCACTAAGGCATCCTTTCCT
GGCTTTTCTAGCCAATCTGGACTCTTTAAGTGTCAGGAGATAACCACTCTCCAAGACAGGCCTTGCCAGTATTCTTGGA
CATCTGCCCTACATGAAAGGCCCTACTTATATCCAGTAGAACTCAGACTCTTTGTTGCACCTTCCATACAACAGATCAC
CCTTTTCCTTGAAGGAAGAGCTCATCATCTTATTCTTTTGCAGGGTCAACAGCCTTAATTTCCTTCACCTTCAGCTTCA
TATAAGCCAATTAGGGTTTTTGTAAGGCTAAGTCAATTCTGTTAACACTGCTTGTCTACATGCATATTTCCTTCCTTAA
ACGAACTCTAATAATTATCCAGTATCTAATAATTATGCCTTCCTTTCAATATATGTTTGTTTATTTGATTCCACTGTAG
AAATACAATAATATGTGCTAAGAACCTATATAGTTTTAAATTTTTCATTTCTATATGCTTACCTATCTGTAGATAAAGG
TTCATAAAGGCATTTATAGACACTATAAAAGTTCACCAGAAACTGCCTTTTAAAAGATAAACACTATTGTTTTATCTAA
AGAAAACAAAAAATAACAAAAAAATACTGTACAAACCTACTCCCTACTAGTCTAAACAGCTCTGCTCCTGTAGTTTGGG
AGCAGAAATTTAAGTGTGCAAATTTGTATTTCTATAGTTCCGATAAAATAATAGAATTTCTCAGTTGAAAATGTCTTAA
GCCTCTCCTCTCCTTCCTAGCTCCATTAAAATAGCCATTGAATAGTATTGGGTATTTCTTTCTTAAAAAAAGATATATA
TTTCCAAATATATTTTCTATGCATTCGTTATATATTTTCTATGCATTCGTTAGAAAAAAGATCAAATACCTGTGCTTTT
AACCTTTTTCTTTTCTTTTCTTTTTCTTTTTGAAACAGAGTCTTGCTCTGTTGCCTAGTTTGGAGTGCAGTGGCACGATA
TAATGGCTCACTGCAGCCTTGAACTCCTGGGCTCAAGCAATCCTCCTACCTCAGCATCCCAAGTAGCTGGGACTACAGG
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TCCTGGGCTTAAGCAATCCTCCTGCCTCGGATTTGCTTTTAGTCTTTACAGTGATTAAGTGGAAAGAGAATAAAGCATC
AAAAGTTTTTTAATGATTATCTTGTAGCTCTGGGTCTGCTATTTACAAGAAGTAGGAGAGAGGCAGAATCTCTCCCTGC
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TATGATTCCTTATTGTTCTTGTACCCTCTTTGAGTACAAACAATCCTGTAACACAGGATCCTCACTTTATATAGTGCAG
GGTAGGTAATTCACAGTTATCAGGAATACAAAACCTTATGCTCTGGATGTGTTATAGATACTAAGAATAATGTCATATT
CTGCTGAGCTCATGGCAACTCTAGAGGAGAGGTTAAAGATCCAATTCTTTCACTTTAGAGAAAGCTGAGACCTACAGAG
TGAAATGACCTGCCACATGTTAGTAGCATTAGAATTCACTCCCCAAGATTCATTCCAGTGTTGTTCCTGCTCTTATGTT
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AACTTGCCTTGATGTACAATGATTCCTCAGTCTTAGAGAACCATCATTTGGCTGTGGGCTTTAAATTGCTTCAGGAAGA
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GGAACTGCATTTCAAAACATATTATGGCCCTTTATGTTATAGAAGCTGCCCGTGATGCTGGCTGTGATGTTCTGTAATA
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TGGTGGCGGGCAGAGAAAAATGACTAACAAAAGCAGATTGTGTGGGCCACAGCTCAAATGGATTTTTTCCCCACCTTTT
CTCATCAGTAGACAGTGCCATTTAGACATCCATGACTTTACTCTTTTTTCTATGCATCTTATTCAGTGATTATGAGACA
CAGGAAAATCTCTAGCTTTCAAAAACTTATAAACTTGTGATGATGTCTTATCCATGGAGATGTCACCCATTTTTCACCA
TGAAAGTGGTTGTCAGTGCCTAGCATTTCTGTATATTACACACATTTATCTGGGCTTTGGGAAAACTTGATAGCAAAGG
GGAAAAGACTCTGCCCCCAAGGAGTAGTAAGGATTTTCCACTGTCATTAAAAGGCATAGTGTTGTTTTATTCCTTTTTC
ATTCTTATATTCTGCGTAATATTTTCATGTGTAAATTCTGTTTTCTCTGAACTTAATAATATACTCTATATTTTAAGGT
ACTTGCAACAGATATGTCAAAACACATGAATCTACTGGCTGATTTGAAGACTATGGTTGAAACTAAGAAAGTGACAAGC
TCTGGAGTTCTTCTTCTTGATAATTATTCCGATAGGATTCAGGTAAAGCCTTGTTTGAGTTTGCTCTGTGTGTGTGTGT
CTGTGCACATGTGCACATGTCTGCGTGTTCAGCTTTGATAAGATGTATTTCTTCCTTCTCCCACCTTATACTTTCAGCA
```

Fig. 6.⌐349⌐

ATTGTAGGCAGGATTTTTATCCAAATTTTTATTCCTGAGAATTAACCAGGTAAAATTCTACTGGTCTTTCTGTTTGCAT
CTACATTAATTAAAAAAACTAACAAACAAACACCCCACAGAACCAGCCACTTAAGCAGCTC. \ATCTAGTCAGCCATG
CACATAAACAGTTTCTCTTAAGCTATTTAGATGCAGTAGAAGTGGCATAATTTGGAACTATTAATACAAGTGTGAACTA
TACACAGACACATCATGGTTGAGCTGTTTGGAATAAATCTTACACTACGTGTATTTTTAAGTGTYGCAGTCATCCAATG
TGAATTTCTAGTTTTTTTTTGTTTTTTTTTTTAAAAAAAAGGAACAGGAGTGAATAGAGAATGCATCCATCTATTTTAGG
TTCTTCAGAATATGGTGCACTGTGCAGATCTGAGCAACCCAACAAAGCCTCTCCAGCTGTACCGCCAGTGGACGGACCG
GATAATGGAGGAGTTCTTCCGCCAAGGAGACCGAGAGAGGGAACGTGGCATGGAGATAAGCCCCATGTGTGACAAGCAC
AATGCTTCCGTGGAAAAATCACAGGTAATGCATGAAGTGTATAGCTTTCAGAGAGAACAGAGCTACCGCTTTAGCATTT
GGTTACTTTGTATTACATATGATAGTATTTTACTGGATTTTTAAAATTACTTTGTTTTTGACAAGCTCAATTTCACCTT
AGTATTTATGATCCAAAGAACTTTCATTCTTATGACTTACATTCATATAGTCATATATATATTCTAAAGACATATTCAT
TTATTATGACTATATTCATTTAATATATTCATATACATAGAGCACATGGCATTATTTCAGTTATCTGGATTCACCTACA
AATTGGTGATTGTAAAATAAGCCCTACCATGTCAACAACTGGAAAATTTTTTATGCTATAGAACATGCTCTTTAACCAA
AGGTTCTAGAAGCTAATTTTGACCAGCTAGTAGCAATACTTTACTTTAAATGGTCTGTTGTTGTTGAAAATAGTGACAA
TTTTACCAAACTAAGTTTAGTAGTCTTCTGTTCAGTGTTTTATT iGTGGGCCATGATCTAATTAAGCTTTTTCCATTGTT
TCTTAGTCCCAAGTCCTCTACTCATACTGGATTTTTTTCTTAACTAGGTGGGGCTTCATAGACTATATTGTTCATCCCCT
CTGGGAGACATGGGCAGACCTCGTCCACCCTGACGCCCAGGATATTTTGGACACTTTGGAGGACAATCGTGAATGGTAC
CAGAGCACAATCCCTCAGAGCCCCTCTCCTGCACCTGATGACCCAGAGGAGGGCCGGCAGGGTCAAACTGAGAAATTCC
AGTTTGAACTAACTTTAGAGGAAGATGGTGAGTCAGACACGGAAAAGGACAGTGGCAGTCAAGTGGAAGAAGACACTAG
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CACTGTCATCTGCCAGGACGTTTGTTGAACAAAACTGACCTTGACTACTCAGTCCAGCGCTCAGGAATATCGTAACCAG
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GCAACAAATATGTCAAGAACAGGACATAGCACGAATCTGTTACCAGTAGGAGGAGGATGAGCCACAGAAATTGCATAAT
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TATGTAAAACAGATTTTTTGTAGAGCTTACTTTTATTATTAAATGTATTGAGGTATTATATTTAAAAAAAAACTATGTTC
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GCTCTAATAATCCACACATTTCATGCTCATTTTTATTGTTTTTACAGCCAGTTATAGTAAGAAAAAGGTTTTTCCCCTT
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GGCTGATTTCAGCACCAGGAACTGTATTACAGTTTTAGAGATTAATTCCTAGTGTTTACCTGATTATAGCAGTTGGCAT
CATGGGGCATTTAATTCTGACTTTATCCCCACGTCAGCCTTAATAAAGTCTTCTTTACCTTCTCTATGAAGACTTTAAA
GCCCAAATAATCATTTTTCACATTGATATTCAAGAATTGAGATAGATAGAAGCCAAAGTGGGTATCTGACAAGTGGAAA
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AGTAAGTCATCAGCCAGGCAAATGACTCATGCTGAAGAGAGTCCCCATTTCAGTCCCCTGAGATCTAGCTGATGCTTAG
ATCCTTTGAAATAAAAATTATGTCTTTATAACTCTGATCTTTTACATAAAGCAGAAGAGGAATCAACTAGTTAATTGCA
AGGTTTCTACTCTGTTTCCTCTGTAAAGATCAGATGGTAATCTTTCAAATAAGAAAAAAATAAAGACGTATGTTTGACC
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AAGAGTCTAGAGTTTATTCCTCTTTCCAAAACATTCTCATTCCTCTCCTCCCTACACTTAGTATTTCCCCCACAGAGTG
CCTAGAATCTTAATAATGAATAAAATAAAAAGCAGCAATATGTCATTAACAAATCCAGACCTGAAAGGGTAAAGGGTTT
ATAACTGCACTAATAAAGAGAGGCTCTTTTTTTTTCTTCCAGTTTGTTGGTTTTTAATGGTACCGTGTTGTAAAGATAC
CCACTAATGGACAATCAAATTGCAGAAAAGGCTCAATATCCAAGAGACAGGGACTAATGCACTGTACAATCTGCTTATC
CTTGCCCTTCTCTCTTGCCAAAGTGTGCTTCAGAAATATATACTGCTTTAAAAAAGAATAAAAGAATATCCTTTTACAA
GTGGCTTTACATTTCCTAAAATGCCATAAGAAAATGCAATATCTGGGTACTGTATGGGGAAAAAAATGTCCAAGTTTGT
GTAAAACCAGTGCATTTCAGCTTGCAAGTTACTGAACACAATAATGCTGTTTTAATTTTGTTTATATCAGTTAAAATT
CACAATAATGTAGATAGAACAAATTACAGACAAGGAAAGAAAAAACTTGAATGAAATGGATTTTACAGAAAGCTTTATG
ATAATTTTTGAATGCATTATTTATTTTTTGTGCCATGCATTTTTTTTTCTCACCAAATGACCTTACCTGTAATACAGTCT

Fig. 6.350

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TGTTTGTCTGTTTACAACCATGTATTTATTGCAATGTACATACTGTAATGTTAATTGTAAATTATCTGTTCTTATTAAA
ACATCATCCCATGATGGGATGGTGTTGATATATTTGGAAACTCTTGGTGAGAGAATGAATGGTGTGTATACATACTCTG
TACATTTTTCTTTTCTCCTGTAATATAGTCTTGTCACCTTAGAGCTTGTTTATGGAAGATTCAAGAAAACTATAAAATA
CTTAAAGATATATAAATTTAAAAAAAACATAGCTGCAGGTCTTTGGTCCCAGGGCTGTGCCTTAACTTTAACCAATATTT
TCTTCTGTTTTGCTGCATTTGAAAGGTAACAGTGGAGCTAGGGCTGGGCATTTTACATCCAGGCTTTTAATTGATTAGA
ATTCTGCCAATAGGTGGATTTTACAAAACCACAGACAACCTCTGAAAGATTCTGAGACCCTTTTGAGACAGAAGCTCTT
AAGTACTTCTTGCCAGGGAGCAGCACTGCATGTGTGATGGTTGTTTGCCATCTGTTGATCAGGAACTACTTCAGCTACT
TGCATTTGATTATTTCCTTTTTTTTTTTTTTTTAACTCGGAAACACAACTGGGGAAATATATTCTTTCCCAGTGATTAT
AAACAATCTTTTTCTTTTTTTTAAGTCCTTTTGGCTTCTAGAGCTCATAGGAAAATGGACTTGATTTGAAATTGGAGCC
AGAGTTTACTCGTGTTGGTTATCTATTCATCAGCTTCCTGACATGTTAAGAGAATACATTAAAGAGAAAATACTGTTTT
TTAATCCTAAAATTTTTCTTCCACTAAGATAAACCAAATGTCCTTACATATATGTAAACCCATCTATTTAAACGCAAAG
GTGGGTTGATGTCAGTTTACATAGCAGAAAGCATTCACTATCCTCTAAGATTTGTTTCTGCAAAACTTTCATTGCTTTA
GAATTTTAAAATTTCACCTTGTACAATGGCCAGCCCCTAAAGCAGGAAACATTTATAATGGATTATATGGAAACATCCT
CCCAGTACTTGCCCAGCCCTTGAATCATGTGGCTTTTCAGTGAAAGGAAAGATTCTTTTTCTAGGAAAAATGAGCCTAT
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TTTTAAACAGTACAAAGGGGACACTATAATTACAAAAACATCCTTAACTGATTTGAGTTGTTTTTATTTCTTTGGATAT
ATTTTCAGAGTGGTAAATTGTGTGTGAGAATTACAAATGATTATTCTTTTAGTGGTTTCTTAGCCTCTCTTACAGCCCA
CGGGGATAGTACTGTACATCAATACCTTCATATGAAATTTTTATATGCAATGAAAATAAAAGCATGGGTTGATTCTGCC
TATTTATGACTCAATCTTTTACAAATAAAAGATTATTCATTTTAAATTATAGTTCAATCAGCATGTCTCTTAGGATACT
GAACGTGGTTGAAATGAAAGGATAGTGACATCATAAGTTAGTACTGATATTCATAACCAAATAAAGCCAACTTGAGTAA
TTTTGCTACATTAAAAATTACCAAAATTACTTAGATGGCCTATAAGATTAAGCATGGTGTTTTCTAAGCAAGCTTTGAA
AGGGGCCTTCCATACTTACTTAATTGAATATTCTGGGATATTGAAAATTATTCAGATACTTGACAATTATTTTTGGTTA
CCTACTCCGCAAACTACAAAGTTTTAAGGACTCAACAATAAGTTAATGAGACACAGTGTTTGCTTTCATGGAGCTTACA
GTCTGGAGGGGACAAAGGCTTAAACAATACTCATATAATTATATATGTGATCAGTACAATGAAGGAGCTCAGTGGGGTA
AATAAGCAGGAACCTGAACTTGATCTGTTCCGGAGGGCCACAGAAGGCTTCCTTGAGGCYTTGAGAAAGTGATTTGCAT
CTGAGTTCTGAAGGATTGTAAGAGGTAACTAGGGAAAAAGTTGACAGGAAGAGGAAGGGGATCCAGACAAGAAACATTT
GCAAAGATCTTGAGGCATAAATGAGCTTGAGACATCTGGAGAAACTGAGGAAAAGTGAGAGAGTAGGCGGGCCTGGAG
CCGCAGAGCCATTGCTAACCATCCTGTGTGAGATATCCCCCATTCTGTAGCTTTATTCTCATAACCCTGCTCAATTTTC
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TAACTAAAAACTCTGTTTGCTTATTCCTCACAAATTCTACTTTTTTCTAAATGACAATCCATTTGTCATGATAATGAGA
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AATAACTGTTTTAAAACTCAAGCTCCCCTTCCCCAAAGCCATGACCCCAAAGTGACACTATGGAACTAAGGAAGCAACTCC
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GACCTGGAAACCTTTCCTCAGGCACACGGCTTTGGACCCCACAACAGCAACATTCCACCAAAAAACAAACCAAAACCAAA
AACATTCCCTGTTGCTAGAGAGTACCACCTGTCTACCCAAGGGGAAAACAACCTTGTGTCAGGGGAATCATACCAGGGCT
TTCCCATCCTTCTGATTTGGGTCCTGCATCCCACCCAGCTACTCCGAGAGCTCCAAATACTCTAGGTCAAACTCAGTGC
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AGGCTTAAAACCAAAAACGAAACAGACTAGAGGTAGTTCTGACTTCCATTTGCTTCTGCTCTCCACTCTGCAAAAACCC
TGGCTATTATTTGAAATATGGAGACGAAAAAGATATTGGAAGAGCATACATTAAATAATAGTCCATTAACCTTCCACA
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GGAAGTCATCAGGAACTCCCATACTTCCTATGTGGGAACACAGGGAAGAGGCATTTTTTCCTGAAGTCTCTGTGTTCC
AGTGCTATCCCTGAATGTCTATTCCCAGCTCTCGCTTAGCTGTTTCAATGACAAGATATAGCACTTGAAAATTTTATAA
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AATACTGAGATTAATGATTTGCAAACTACAGATTTGACAGAAGTCCTAATGCTATCTCTACATCCTATTTCTGTTGAGG
ACCAAAATTGACTGTTGCATTGCCTTTGATTTCCTTTGACCATCCTTCATTACAGATTTCAGTAATTTTTATGGTCCAA
AGCTAGGTGACAGATATTACATCATATGCATTTGTTAACTCACCCTATGTCTGCATAGCCTTGCTATATGGTCAGAATT

Fig. 6.351

CTCCAGATTGACTGGGGAACTTGGAGTTAAAGAGCCCCTGACCCCCTAAAAAATTATAGAAAATGAAAGAAGCTGTCAA
GAAAGTAAGGAGCAGAGATAAGAGCAGAGATGGGAGYGTGCCTCTGCCCGGCAGAGACTCACAACATGAGGCAAGACAG
ATCTTAGCTGAACTTCTTGCAGGATGTATGCCTCAATGAAAGGAATGGAGCACGGGCAGCATCCGGCTTAGGCTGGAGC
CCCTTCAACCATCATGGCTGTTTTAGCAGGGTGTTTGCCAGCTGCTTACAAATCCCAAGCCAACAGACATGGTTTCAGG
ACCTGAATCATGCAAACATATTTTCTGAAAGGTTCTATCACCTGAGAATTGCTTTTCCTTCGCTCCTTGTTTTTGAACC
AAAAAGCCGACCAAATGTATAGTGGCAAACACTTGCTCAGCACCCATCCATGCAAGGCACTGTTTTACCACCAGGCTGC
ATCAGTTAGAGAACCAAGCCGTCTCTCACCTTGCAGCTTCTTCACACTTTGCTCATTTTCTCTCTAACACTGATGGTTT
CTGTTTCCTGCTATTCAGTTACAGCATCTGGGAAGAGAGTGGCAGAGAACGCCAGGCAACTGCACCCTGCCACAGTGGG
TCAAGCGGAGGCTGGGTGGTGACACCCGTCTCTACCCAGCAGAAGTCAGTGGTTCATGGGTGGAGAACAAAGCTCAGGA
CAGCTGCTGCAGTCCAGGGAACTGTCCACCATAGAAGACTCTTGGGAAGCAGCCCCTGACCCCAACCCCCTGTGCTTTT
TGAAGCCGTGTCTGGTATCACATGGAATTATGCAGTAGTCATAGAATATCTTTGAAAACTTATATTCTAAAAATAATTG
CTGAGATCAGCGATTAAAAATATCTAATAAATGGACTGGAAAAGTTGAAAACAAGCAAAGGTGAGAGAAAACAATTGAGG
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CAGGACCCAGGTTCCTGGGTGCTTAACGTAACACATAAGGCCCTTAAGGATCTAATCAAACTAAATCATCCCCAACTCC
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GAAAACAGCTCATCCCTCTTGATCCAGTTTAAATACACCTTTGTCTGGAAAGCCTTTCCCATTAGTATTTTATACACAA
CCCTATAATAAATATTTCCCCTAATATGTTGTATTTTTTAAAAAAATTCTTTCTCTCACTAGATTGTGAGATTTTTAAT
TTTTATTTTTTTAGACAGTCTCACTCTGTCACCTAGGCTGAAATACAGTGGCGCAATGTTGGCTCACTGCAATCTCTGC
CTCCTGAGTTCAAGTGATTCTCCTGCCTCAGCCTCCTGAGTAGCTGGGATTACAGGCATGCATCACCACGTCCAGCTAA
TTTTTGTAATTTTTTAGTAGAGATGGGGTTTCGCTATGTTGGCCAGGCTGGTCTCGAATCCTGACCTCAGGTGATCCGC
CCACCTTGGCCTACCAAAGTGCTAGGATTACAGGCGTGAGCSACCACGCCTGGCCTAGATAGTGCAATTTTTAAACATC
AATGTATTCCTTTTTGCCTGCTAGTAAGGAATTAAGTCCGGATATTAACATTTGACTATTCTTTCCCCTCTAACACCAA
CTCTTCTCAAAATGTGGTCCTTGAGCTGACATTAACGTCAACTAGGATCTTGTTAGATTATAATCTAGCCCCAGACCAA
GGCTCCAGGAAAAGACTCTCTAGGGGTGGACCCTAGAACCTGTGTTTTAACAAGGACTCCAGGTGATAAGTATGCTCCA
TAAAGTTTGAGAAACACTGTTCAAGACATCATGGAGGTGTTTTTCTATTTCTGTTTTTTTTAAGAAGCAAGTTAATAAT
CTCAACCACCTTTCTAATGTATTTCATTTAATCAAAGGGATACTTGTTGCTACAGGAGTTACAACAGATGACCTAGTGA
GGGCCACAGGCAGTCTGAGGAACAGTAGGATCCCATGTTTGGCCAATGTCTATCAAGCAAGAAAGGAATAGTGAAGCCA
GCTATTAGCTCCCTTCAGCAGAGGAAACCTGGGATTAGGAGTCGCCATCAGGCATCATGGCATCACAAGCAGGCATGGG
TTRAAGATTTAAATGTTGCAAAAAGAGAGAAGAACCAGAAGGAAATTACGTCCACATTATCCCTGCAAGACCATTAATG
TAATGAACAGAAATCAAACACCTTTTCAGTCTTCAAGTGCCGCATTCAGGTTCACTCTGCTAATCACACAGCTGAGTGG
CCCTCTGCTTTCATAACTGGGTGAAACCACACATCCCACCCCAAGATGGCTTGGGATCTACCATCACAAGGGGCCACCA
AAGGGCCAAATAACACAACCCCGAAATTCAGTGGTGGTAACTCTCCATGGGGTTAACTTTGATTACTGGGAAACTAGAA
ATGGGAAGGAGCCAGTAAATTAATTTCTTCTCTCCCAAGACTATTCTGAGGCATAGCCTCCCCCTTGAAGCCCATCAGG
ATGATCCTGCAGTTCCATGTCAGCAAGAGCGCCCTACCAAGTGACCTGCAGTGTCTCTTGGCTGCTCCTTGTGAGTCAG
TAGCCAGCAGAGAAATGTTTTCCCTACTTTCCTTTATTTCTTTGTTCTTACTTGCTTCATTTTGCCTTGACCTTTTTTT
CCTCACCCCTACTACCCAAAGGCATGTACTTCCCAAATAAAGTCTGTCACTCTGATCCCTCCCTCAGACTGTTTCCAGG
GATGCTGAGCACCCTGTACATTCAGAAAAGCAGATGTCAAATTTCTGTATATATATTGGAGTGCACTAAGTATTTAAAA
TCAACCTTTGGCTATTAGAGAGGTAGGGACTGTGCCTTATGCATCATTGTTTTTCCAGGGCTTAGTTCAGTGTCTGCCATA
TAGCAAGTATTCATTTAACAGGAAAAGAGAAAGACAAGTGGGTAAGTAATTGCCTGAATGAATGAGGTTGATGGTGAAG
AATGGGTGGGAAGGGCATGAATTGGCCTAGAATAGATTAGACATTAAGTAAGCAGGAGTCGAGACAGTTAAGGGGCCAA
AGGCAGCTGAAAAGAATATGGAACAGGAATACAAGTTGAAGGAATCAGGCATTCTCCCTAGAAACAAAGGGTACGGGAA
GGGTAAGAGACAGACACTTCACTGTCTGGGAATTTTCATCCATACTCGTTATTAAAGACACACTTGGTTGCCTACTTAA
TTATTACCAACCCTCCACCCCCACTTTATTCCTGGCATACATAAGTGGGATTTTGTTCAGGTCGTGTGAAGTCAAGTCT
TCCAGGGAGATCCAGCCTCTCCTCAGTTCTGTAGGCTAATTCTGAGCCTTTTATGGTAATTCCTCTACCCTTGCCAGTG
ATTAGGTTAGGACAGGCTTGCCGTGCTGCAATAGTAGACAGTGAGACACAGGTTGACTTCTGGGAAAATATCCCCAACT
CTTACAAAGAAAGAGAGAACAGAACATTTTTTCTTCGGGGCTTTGGAAATTATTTTCAAAGATGTCATGGTTAGGCTA
TTATCTTGAGGCCACGAAAGGAACTATCTGAATACACAGACAACATACTGAAAATGGCAAAGCAGAAGTTAGATAGCAC
ATGAGTCTTTGAAGATGTGTTTGAGCTGCTGAACTAATCAGCCCTACTTTGAAACTTATTAAGATAATAACTAGTTATT
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CGTTATAAAGGAATACCTGATGCTGGGTAATTTATAAAGAAAGGAGGTTTATTTGGCTCAGAGTTCTTCAGGCTGTACA
GGAAGCATGCACGGCTCTACAGCTGCCTGGCTTCTGGTGAGGCCTCAGGAAGCTTTAACTCATGGCAGAAACAAGGGGA
GCTGGCGTATCACATAGTGGGAGGAGGTGGCAAGTAGGCAGGGAGGAGGTGCCATGCTGTTAAACAACCAGCTCCTTCA
TGAATAGAGTGAAAACTTATACATTACTGCAAGGACAGCACCAAGTGGATCATGAAGGATCGTGACCCAAACACTTCCC
AATAGGCCCCACCTTGAATATTGGGGGATCAATTTCAACATGAGACTTGGAGAATACAAATATCCAAACTATATCATAC
ACCCACTCTTTGTTCCAACAATGTGCTGGTGAGTCTGGGCTGGGARTGGGGGAGAAAGGTACCTGGCACAGTGGAAATG
TTGGAAATTTGACCCCAAATTAGCTGGGTGACCTTAGGCATGTTAATTCATCTCTACTTTCTTTCATTTTAAATAGGGG
AATTATAATATTGGAATCTTCTCTACATTCTTCACAGTATTAGAGCAAATGAATTTATAAATGTAAACTCAGTGTAAAC
TTTAGATAAAATTTTTCAAACACTAAGCTTGTGACCCAGAGTAGATAATAAAGTCAAAAGATGGTTGCCATGAGCACTA
ACCGTGAATTAAAAATAGAATAGAGAATATCACTGTGTGCTGCACATAAGTTTCATTTTCTGAAATTGCTTCATTTACAT
GCATGCATGGGTGCACTTGTGTGCTTGCACACACATGCAAATGTTGCCAAGTAAACTATTTTTAAAAGCATACGAAGCA

Fig. 6.352

TGGATTTAGAACAAAGTCAACACCAGAGACAACTAGAGGAACCTAGTCTCTCACCAACCCCCACCCCTGCCAATTTAC
CTGTCATCCATGATTCAAAAATGGTTTCTAAAGGGAAATAAAAATTGATTCAAAAGGGAGACCTAATGAGTAACATAAT
AGGGAGCCATTGATTGTTTTAGAGTGGTATGACGGAGCAGTGGGTAAGTAGATCACTCAACACTGGTATTTCAGACTGC
CAAGAGAGGAGAAAACAGACACAAAGTGATTTCTGGGCTTTTGACAACTTTTTTTTTCTCAAAGTAGATTGACAATATT
ATGTGTCATCTTATATATATTAATAGACAAAGGCAATATCTACACTTGGGAATACTTCATGGAAGGCTTGGCTGGAAAC
TAAAAACTGGACTATAAACTTCTTGGGGGTTAGATACGTATCTTAAGACTTCCTTGTGTCTTGAATTGTGCCTAGCCTA
GTGCTGCATAGATATGTAAGCCACTCATTCTTGTGCAATTACCTATTGTAGAATTTTCAGACTAATATATTTCCTTTTC
CTTGTTTTGATCCTCTGCTGAATCATGAGAATAGTTCTCAGGTGAACAGCAGTGATGTACAGTGTCTCACTGGAAAAGC
AGGAAAATAATCAGATGTACCCCAGAACCTTAGGATATGGGAGGCTCTTCTACTGGCCACTCCATGGGAAACTCGCTTG
CTTTGGGGGACAGTTTGCTTTGCTCTATTTTGTGGGCACAGGGTTTTGCCTTATTTTTCTACTCCCTTATTCCCAGCAA
ACATCCTAAAAAGATGTGGGTAAAAGAAAAAGAGAATAAATTTTGCAATTCTACTCACTGGCTAGAAACAGACAGAGAC
ACATGAGTAGAGGCGGAGTTTACCATCACCCTTAATCATCCTGGGGCCCATGTGTGTATAAAAGGCAAGAAAAGAGCCA
TTACCAGGAGGCACTCACCTTCACAGTTTCCACCGCATAATTCCACCCCTCTGTATCAAACTACCCAAATTGCTACACC
AATCACAACTGCCCTCTTGTTCAAAGAAAATAATATCATGGATTTCTTTCCTGGCAGTTCTGCTATTGATTGTTTTTGT
TCTTTTAATAAGCAAGGTGAGAAGGTTAGAAAAAAATTATTCAGATAATTCACCTGGGAGTAGGGTAGGAACTTGAGGC
ATGCAGAGAGAATGGCAAATTCAAAATCAACTAAGCCATAACTGCCTATCCTACTGACCACTGTGCCAGGTACCTCAAA
AGAGTAGGACTTGATTCTTCCTTTTGAGGAGGTTTCAGTTTGGTTGGAAAGGAATGGGGGAGGGGCCAAGTACAGTGGC
TCACACCTGTAATAGCAGCACTTTGGGAGGCCGAGGTGGGTGAATTGCTTGAGCCCAGGCATTTGAAACCAGCCTGGGC
AACATGGCAAGACCCCACATCTACAAAAAATACAAAAATTAGCTAAGCATGATGGTGTGCACCTGTAGTCCCAGCTACT
CGGGAGGATGAGGTGGGAGAATCACTTGAGCCCAGGAGGTTGAGGCTACAGTGAGCTGTGAACATGCTTCTGTGCTCCA
GTCTGGACTACAGAGCAAGACTCTGCCTCAGATAATAAAAAAATAAAAAAGGAAAAAAAAAAAAAGAAATGGGGAATGGA
AGAGGAAGGAATCCAGGGGAAGAATGATCACTAAGACTGCATCACACCTTTTGCTATCTCATTTCAACTCTACATCAAC
CCAATATTCCCTTCATTTAACAGAGAGGTCAAAAGAGGCTGGAAGGATAAGGTTGTCCAGTAAAAATGTCAAGGCTGAT
ATGGGAACATAGCCAGTTTGTCTCTAAAGTGCCCTGTGTCCCTTGGGGAAGAGAATATTTAACTTGATTGTTGCTTCAG
TTTTTTATTCTATGTCAGCTATCAGTTTGCAAGGCATTATTTAGGGGATTTCCTTTACTTTCTTTTTTTTTAAATCTCA
AATTTCCCTCTCATTTAGTAACTGTAATTTCAGTGCCTAACATGTAGGACAAAATACTTTCCATATTTTTTTTTCAAAA
TTTTTGCACTGACTGTAGTCCCCTCACTGGAACAGCTTTATTTCCCTAAATAATATACAATGAACTTGTTCATATCGAA
GACGTATGCCAATATTAAATACAAACAGCTCAGCTGGGCGTGGTGGCTCATGCCTGTAATCCCAGCACTTTGGGAGACT
GAGGCAGGTGGATCACCTGAGGTCAGGAGTTCAAGACCAGCCTGGCTAACATGGAGAAACCCTGTCTCTAATAAAAATA
CAAAAATTACCCAGGTGTGGTGGCACATGCCTGTAATCCCAGCTACTCAGGAGGCTGAGGCAGGAGAATCACTTGAACC
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GCAACCCACAGACCAGGAAGTTCCCTTGGGTGACTATGCCACCAGGGCCCTGGGTATCAAGCACAAAACTGGGCAGCTG
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TCATTCCCCTGGAAAGGGGGCTGAAGCCAGGGATCCAAGTGGTCTAGCTCAGCGGACCCCAACCCCACAGAGCCCAGCA
AGCTAAGATTCACTGGCTTGAAATTCTCACTGCCACCACAGCAGTCTGAAGTCAACCTGGGGCACTCGGGCTTGGTGGG
GGGAGGGGTGTCTGCCATTACTGAAGCTTGAGTAGACTGTTTTCCCCTCACAGTGTAAACAAAGCCAAGGGGAAGTTCC
AACTTGGTGGATCCCTCCGCAGCTCAGCAAAGCCATTGAAGCCAGACTGCCTCTCTAGATTGCTCCTCTCTGAGCAGGA
CATCTCTGAAAAAAAAGGCAGCAGCCCCAGTCAGGGACTTATAGATAAAAACCCCCATCTCCCTGGGACAGAGCACCCA
GGGGAAGCAGCGGTGGGCACAGCTTCAGCGAGACTTAAACATCCCTGCCTGCTGGGTCTGAAGAGAGCAGCAGATCTCCC
AGGACAGCGTTCAAGCTCTGTTAAGGGTCAGACTGCCTCCTCAAGTGGGTCCCTAACCCCCATTGTAGCCTGACTGGGA
GACACCACCCAGCAAGGGTTGACAGACACCTCATAGAGGAGACCTCTCGCTGGCATCTGGCGGGTGACCCTCAGGGACA
AAGCTTCCAGAGGAAGGAGCAGGCAGCAATTTTTGCTGTTCTGCAGCCTCCGCTGGTGATATAGGTAAACAGGGTCTGG
AGTGGGCCTCCAGCAAACTCCAGCAGACCTGCAGAAGAGGGCCTGACTGTTAGAAGCAAAACTAACAAACAGAAAGGA
ACAGCAATGACATCAACCAAAAGGATGTCCACACAAAAACTCCATTCGAAGCTTACCAACATCAAAGACCCAAGGTAGA
TAAATCCATGAAGATGAGAAAAAAATCAATGCACAAAGGCTGAAAATTCCAAAAACCAGAATGCCTCTTCTCCTCCAAA
GGAACAGAACTCCTCAACAGCAAGGGAACAAAACTGGATGGAGAATGAGTTTGACGAATTGATAGAAGTAGGCTTTAGA
AGGTGGGTAATAACAAACTCCTCTGAGCTAAAGGAGCATGTTCTAACCCAATGCAAGGAAGCCAAGAACGTTGAAAAAA
GGTTAGATGAATTGCTAACTGGAATAACCAGTTTAAAGAAGAACATAAATGACCTGATGGAGCTGAAAAACACAGCATG
GGAACTTTGTGAAGCATATGCAAGTATCAATAGCCAAATAGATCAAACCAAAGAAAGGATATCAGAGATTGAAGATCAA
CTTACTGAAATAAAGCATGAAGACAAGATTAGAGAAAAAGAAGGAAAGGAAACAAAGCCTCCAAGAAATATGAGACTA
TGCGAAAAGAACAAACCTACATTTGACTGGTGTACCTAAAAGTGATGGGGAGAATGGAACCAAAAGTTGGAAAACACTC
TTCAGGATATTATCCAAGAGAACTTCCACAACCTAGCAAGTCAGGCCAACATTCAAATTCAGGAAATTCAGAGAACACC
ACAAAGATACTCCTTGAGAAGAGCAACCCTAAGACACATAATCGTCACATTCACCAATGTTGAAATGAAGAAAAAAATG
TTAAGGGCAACCAGAGAGAAAGGTTAGGTTACCCACAAAGGAAAGCCCATCAGACTAACAGTGGATCTCTCTGCAGAAA
CCCTACAAGACAGAAGAGAGTGGGGGCCAATATTCAACTTTCTTAAAGAAAAGAATTTTCAACCCAGAATTTCATATCC
AGCCAAACTAAGCTTCAAAAGTGAAGAAGAAATAAAATCCTTTACAGACAAGCAAATGCTGAGAGATTTTGTCACCACC

Fig. 6.353

AGGCTTGCCTTACAAGAGCTCCTAAAGGAAGCACTAAACATGGAAAGGAAAAACCAGTACCAGCCACTGCAAAAACATA
CCAAATTGTAAAGACCATCAACACTATGAAGAAACTTCATCAACAGGCAAAATAACCAGCTAGCATCATAATGACAGAA
TCAAATTCACACATAACAATATTAACCTTACATGTAAATGGACTAAATGCCCCAATTAAAAGAAACAGACTGGCAAATT
AGATAGAGTCAAGAAGCAACGGTGTGCTGTATTCAGGAGACCGATCTCACGTGCAAAGACACACATAGACTCAAAATAA
AGGATTGGAGGAATATTTACCAAGTAAATGGAAAGCAAAAAAAGCAGGGGTTGCAATCCTAATATCTGATAAAACAGAC
TTTAAACCAACAAAGATCAAAAAAGACAAAGAAGGGCTTTACATAATGGTAAAGGGATTGATGCAACAAAAAGAGCTAA
CTATGCTAACTCTCCTAAATATATATGCACCCAATACAGGACCACCCAGATTCAAAAGGCAAGTTATTAGAGACCCACA
AAGAGACTTCAACTCCCATACAATAATAGTGGGAGACTTTAACGACCCACTGTCAATATTAGACAGATCAATGGGACAG
AAAATTAATAAGGATATTCAGGACTTGAACTCAGTTCTGGACCAAGCAGACCTAATAGACATCTACAGAACTCTCCACC
CCAAATTCATAGAATATACATTCTTCTCAGCACCACATCACACTTATYCTAAAATTGACCAACAGAATTGGAAGTAAAA
TACTCCTCAGCAAATGCAAAAGAATGGAAATCATAACAAACAGTATCTCAAACCACAGTGCATCAAATTAGAACTCAGG
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AAATTAAGGCAGAAAATAAGTTCTTTGAAACCAATGAGAACAAAGGCACAACATACCAGAATCTCTGGTACATACCCAA
AATAGTGTTTAAAGGGAAATTTATAGCACTGAATGGCCACAAGAGAAAGCAGGAAAGATCTAAAATCGACACCCTAACA
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CAGAACTGAAGTAAACGGAGACACGAAAAATCCTTCAAAAAAAAAAAATCAATGAACCCACGAGCTGGTTTTTTGAAAA
GATCAACAAAATAGACCACTAGCCAGACTAATAAAGAAGAAAACAGAGAAGAATCAAATAGACACAATAAAAAATGATA
AAGGGGATATCACCACTGGTCCCACAGAAATATAAACTACCATCAGAGAATACTATAAACACCTCTCTGCAAATAAATT
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CCTGAAGAGACCAATAACAAGTTCTGAAATTGAGACAGCAATTAATAGCCTACCAACCAGAAAAAGTCCAGGACCAAAC
AGATTCACAACAGAATTCTACTTGAGGTACAAAGAGCTGGTACCATTCCTTCTGAAACTATTCCAAACAATAGAAAAAG
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CAGCACATCAAAAACCTTATCCACCATGATCAAGTAGGCTTCATCTCTGGGATGCAAGGCTAGTTCAACATATGCAAAT
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CGATAAAATTCAACACCCCTTCACGCTAAAAACTCTCAATAAACTAGGTATTGATGGAAGGTATCTCAAAATAATAAGA
GCTATTTATGACAAAACCCACAGGCCAATGTCATACTGAATGGGCAAAAGCTGGAAGCTTTCCCTTTGAAAACCAGAACAA
GACAAGGATACCCTCTCTCTCTATTCCTATTCAACACAGTATTGGAAGTTCTGGCCAGGGCAATCAGGCAAGAGAAAGA
AATAAAGGGTATTCAGATAGGAAGAGAGGAAGTCATATTGTCTCTGTTTGCAGATGACATGATTGTATATTTAGAAAAC
TCATCATCTCAGCCCAAAATCTCCTTAAGCTGATAAGCAACTTCAGCAAAGTCTCAGGATACAAAATCAATGTGCAAAA
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AATAAGAGAGGACACAAATGGAAAAGCATTCCATRCTCATGGATAGGACGAATCAATATCATGAAAATGGCAAAATGGC
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GGCATCACATTACCTGACTTCAAACTATACTACAAGGCTACAGTAACAAATATAGCATGATACTGGTACCAAAACAGAG
ATATAGACCAATGGAACAGAACAGAGGCCTCAGAAATCACACCACCCATCTACAACCATCTGATCTTTCACAAACCTGA
GAAAAACAAGCAATGGGGAAAGGATTCCCTATTTAATAAATGGTGTTAGGAAAACTGGCTAGTCATATGCAGAAAACTG
AAAATGGACACCTTCCTTATACCTTATGCAAAAATTAACTCAGGATGGATTAAAGACTTAAATGTAAGACCTAAAACCA
TAAAAACTCTAGAAGAAAACCTAGGCAATACCATTCAGGACATAGGCATGGGCAAAGATTTCATGACTAAAACACCAAA
AGCAATGGCAACAAAAGCCAAAATTTACAAATGGGATCTAATTAAACTAAAGAGCTTCTGCACAGCAAAAGAAACTATC
ATCAGAGTGAACAGGCAACCTACAGTGGGAGAAAGTTTTGCAATCTATTCACCTGACAAAGGGCTACTATCCAGAATC
TACAAAGAAATTAAACAAGTTTGCAAGGTAAAAAACAACCTCATCAAAAGTGGGTGAAGGATAAAAACAGACACTTCT
CAAAGAAGACATTTATGGAGCCAACAAACATATGAAAAAAAGCTCTTCATCACTGGTCATTAGAGAAATGCAAATCAAA
ACCACAACGAGATACCATCTCATGCCAGTTAGAATAATGATCATTAAAAAGTCAGGAAACAACAGATGCTGGAGAGGAT
GTGGAGAAACAGGAACACTTTTACACTGTTGGTGGGAGTGTAAATTAGTTAAACCATTGTGCAAGACAGTGTGGCAATT
CCTCAAGGATCTAGAACAAGAAATACCATTTGACCCAGCAATCCCATAACTGGGTATATACCCAAAGGATTATAAATCA
TTCAACTATAAAGACACATGCACACGTATGTTTATTGCAGCACTGTTCACAATAGCAAAGACTTGGAACCAACACAAAT
GCCCACCAAGGATAGACTGGATAAAGAAAATGTGGCACATATACACCATGGAATACTATGCAGCCATAAAAAGGATGAG
TTCATGTCCTTTGCAGGGACATGGATGAAGCTGGAAACCATCATTCTCAGCAAACACAAGAACAGAAAACCAAACACTG
CATATTCTCACTCATATGTGGGAGTTGAACAATGAGAACACATGGACACAGGGAGGGGAACATCACACACTGGGGACTG
TCGAGGGGTGGGGGGCTGTGGGAGGGATAGCATTAAGGAGAAATACCTAATATAGATGATGGGTTGATGTGTGTAGCAA
ACCACCATGGCATGTGTATACCTATGTAACAAACCTGCACGTTCTGCACATGTATCTCAGAACTTAAAGTATAATAAAA
ACTCAAACAGCTCTACATTGTTATTTATTTAAAACTTCAATTTACTTGCTAAGAATTATCTTTTTTTTTTCATTCTCTC
TTCACTTCTCTCACCAGCAATGGCATTTGGGTTTTTTTAAGGCCACTTTAAAAGACATTGAAAATAATACCTGTTTGTT
TTCATTCTATTATTCTTATTACATTCATTTGTCTTTGAATGTTCCCAAGTTTTCTGGTATGACACTACGAATCTAAGTT
ATTCCAGACTTCTCTATTCTTTCATGTATTTAGAATACATTTTTCAAAATTCCTAGGCTGAGGTATTAATAACTTGCCC
AAATTACCTTTCAAATGTATTTACCATCCCTGTATTACTCAGTACAAAAATTTGATTTTTTGGAGACATATTTGTACA
TATTTATGGGATACATGTAGTATTTTGTTACATGCACAGAACATGTAATGATCAAGTCAGGCTATTTGGGCTATTCATC
ACCTCCATTATTGATTATACCTATATGTTGAGAACATCTTAAGTCCTCTTTTATAGTAAGTTTGAAACATATAATACTA

Fig. 6 354

TACTATAGTCACCCTACTCTGTTATTGCATATTAGAATTTTTTCCTTCTGTGTGTTTGTACCCATTAACCAACCTCTAC
TTCATTCACCCCCCACCACCCACACACCCTTCCAATCCTCTGGTGTCTATCATTCTATTCTCTACTTCCATAAGATCCA
CTTTTTTAGCTCTTACATATGAGTGAGAACATGTGATATTCTTCTTTCTGTGACTGGCTTATGTCACTTAAGATAATGA
CCCTTCAGTTCTATCCAAGTTGCTGTAAATGCCATCATGTTAGTCTTGTTTATGGCTGAATAGTATTCCATTGTGTATA
TATATATATTCTTTAACCATTCATCCATGATGGATACTTACGTTGATTCCTTATCTTTGCAATTGTGAATGGTGCTGCA
ATAAACATGGGCTGCAGGTATTCCTTTGATATATTAATTTCCTTTCCTTTGGATAAATACTAGTTAGATTACTGGACTG
TATGGTAGCTGTTTTTAGTTTTTTGAGAAATCTCCATACTGTTTTCCAAAATGGCTGTACTAGTTTACATTCCCACCAA
CTCCCACCAACAAGAATTCCCTTTTTTCTGCATCCTCACCAGCATCTATATTTCGTCTTCATCTTCTCCTTCTCCTTCT
TTGACAGGGTCTTGCTCTGTTGCCTGTTTCCCAGGCTGAAGAACAGTGGTGATTCACAGGTGCAATCAGGATGCACTGC
AGCCTTGAACTCCTAGCCTTAAGCAATACTCCTGCCTCAGCCTCATGCGTAGCTGAGACTACAGGCATCAGACTTTTGT
CTTTTTAGTAATAGTCATTTTAACTGGAGTATGATGCTATCTCATTGTGATTTTAATTTGCAGTTTCCCGATGATTACT
GATGTTGAGCATTTTTTAATATGCCCATTTGTCTTTTTTTGAGAAATATCTATTCATGTCACGCAACTTCTGCCTCCTAGGC
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CCCACAGTGCTGGGATTACAGGTGTAAGCCACTGCACCTGGCCCTTTACCCACTTTTTAGTAGGATGATTTGTGGTCTT
TTACTGTTGAGTTGTTTGAGTTCCTTGTATATTCTGGATACTAGTCCCATTTGTCT
CCCATTCAACAAGCTGTATCTTCAGTCTGTTGGTTTCTTGTGTGTAGAAGAATTTTTAGTTTAATATAGTCCCATTTGTCT
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CTATGTTTTCTTGTAGTAGTTCATAATTTTGGGTCTTATGTTTAAGTCCTTAACTGATTTGAGTTGATTTTTGTACAGG
GTGAGAGATGGGTCCAGTTTCATTCTTCTGCATATGGATATCCAGTTTTTTCTATTCCATTTAGTGAAGAAAGTGTCCT
TTCCTCAGTGTATATACTTGGCACCTTTATAGAAAATCAGTTGGTGGTAAATGTGGTATATGCTGGCATCAGTGTTAGT
GTGTCCAGGCGGGCTGATCTGGGGGCTTCCAGTGGTGTGGGTGATGGCAGTTGCAGTGGCAGGACAATCCTCTGGTACCCA
GGCAGGTCCATAGGCCCTGGGCATCAGGCATGGCAGGCCATTTCCCAGGCCCACAGGTGGTTCATGTCTGTGGGTGGGGGGT
AGTAGTCCCAACTGATTTTCAAGGTGGCTGCAGGTGGGTGAACCCATCTCCAGGCCTTCAGGATGAGTGCTCAGGCGCCAACAG
GTATGCTGGCTGTGATGGTAGTGGCAGGTTGGGTGAACCCATCTCCAGGCCTTCAGGATGAGTGCTCAGGCGCCAACAG
GAATAGATGGGGCTGAGCAATCCCCAGGCCCCCTGCATGGGCACTAGGGAGAAGGGAGACAGAGGTGAGCCTCAGGCCCC
CCGATGGTATATATAGGCACTAGCTATGGTAGGCAGGGGCATGGTGATTTCCAGGCCCTCAGTGGAATGCTTGGATGGG
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ACACTTTGGCCCCAGGTGGTAGTTGCTGGTGGGGTAGCCTGTTCTCAGGGTGCTTCTAAATGTACGGTACCCTGCTGAT
GGGGATGGTGGGTAGCTTCCAGTGGCCCCCACATTGGATATGGAGGCAGCAGCCAGCAGCAGCAGGGTCTGCGTTAGGG
GGAGGTCAATGGGGCTCAAGGAATCTGGAGTTGCAAGGTCTGTGGGGTCCCAGGGTAGGATGCAGTCTGCTGGGCTTTC
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CTGAGTTTTGACCAAGCCTCTGATATGCATACACATATATACATGTTCATGTACGTGTACTTAAAACATTTTTATTCCC
ATGTTTAAGGCTAGTTGAATAGGGGTGAACACAGAAAATCCTAGCAATTCATTCTTTGCACTCCTGGCATTCTCAAATG
TTTTGTAGATGCCTTCTTGTAGCAGCCAGTCTCGTCCCTTTTCTTGAAGTCAATGAGCATCAAATGTGTTCAGGATTCA
TGAAATAATTCTGACAATAATTAGTATTTTTTATAGGAATTGGAACATGGTTTATTGCAGTATACCCCAGTGTTTGTCA
GCTCTAGAACTTACGAAGAAGAGGAAGAGGAGAACAAGTTTGACTCCTGTGAACATTTTCCTTTTTTCTATAATTTACG
CATATGCTAACAAGCAGCTAGTTTCAAAACACAAACTTACCTTGACAGGAACTAGGTCTCACTGTTGAGAATGTAGTCT
CCTCTCTCCCCACATAGCCCTAGGGTTATTCAGGATGAAGGAGAAGAAAAAACATACCCAAGAGCATTTTATAATTCCA
TCCTTCCTTCTTGTTATGCTAAGGTTACTAGTACCATGACCTATGTGAACTCGTTTTCTTGAATAAGAAGAATAAAAGC
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ATCATTCCCAGGCCTGCTTTTGATCTGAGCAGTGGCTTTCAAAAAATATGCTGATGAGATTGCTCTCTCACTTAAGAAC
TTAAGAAAGAGTTGAAGTTCTTCAATGTATATCCCCAGCCTGAAACTTGTCTTGAGGAAGATTTGTAGTAGTAATGAAG
GGTGACAGTAACCTAAGGGTGAAGAATGTAGAGAGATTTTTCCCAAGAGCAATCCAAGATACTGCTTGAAACATCCGGG
GTCCACTGACATGCCCCTTGGACTCCAGGAGAAATCTGGAAATGTCTGGACCTGCAATGAGACACACCCAGCTTCAAAT
GTGTCACTCTGGGCATATTCTTTAATCTGAGCTCCAATTTTCTTTCTTGTAAGAGTCATTACACATGGCTGTTTATTT
ACTAATTCGTTTTTTAAATTAAAGAAATAAATTTGTAGGAGTAATATAGTTACATTGTTTACAAGTCAAATTCTTACAT
GGCTCATTTCCATCCATTTCATGGATGTGTTACCTTCTGGTATTTATTAGAATTTGCTTTAGACTTTCTGTTGCTCACC
TCACAGTTCTTGTGTCATCTTAGTTTCTTTTTCTGTTTTTTGGTCTTTTATGCGAGAGGTTTTCCTCACATTGTCTGATA
ATCCTTGGCAGTCAGTTCGTATTGAAGAGTTAGACATTAGAAAGCCAAGTGAAAATCCTGTGTGTGTACAAGGTTTGTT
GACTGGAAAGCTTCATGGGGATATTCTTGCTTGGACATTTCACTGAAGGACATCCAAAAACCTGTTGCCGTTTTTAGAT

Fig. 6.355

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TTTTTTTCTTGGATTGTTGGTTTTCCTGGTGATCTTACTCCAATCTCTTTGGTGCCATAATTCTACCAGCTGAGTATGA
GAAGGAGCTGGGGTTTCTCAATGCAGAATATGTACTTGGTTCTAATATGGTAGTCTGGTTCTAATATTGTAGCCTTTTT
CTCAGCAATGGTTGCCACATCTGAGTCCTGAATCCCTTTGGAATCAGCCTCTCCAGAGGGTGTATGTTTAGAACCGGCG
TGAGTGAAAGGTAGTCATCCAATGACATGAGGTGGGTGAAAATATTTTAACTGCTTTATTTATAAATTTTCAACTGACC
CTCCTGTTTTTGAAACCCACCAGTGTCCCCACTTCCATAAGTCCCTGGATCCTCAATTTTCCAAGTCTTCCTGGAATTC
TATGGAGTAAATTAGCTTACTGATGCATTTCCCTTACTTGCCTTAGGTTTCTGCTCACTCTGTTCCCAAGTCAATTATG
ACCTACCTGGCTGCTTCCAGTTTCTGAAATGTTGTGTGTCAGCTTCTTTTCCATTGCCTTTGTCCTTGCAAGCTTATGC
CATTTTAAAAAATTCTTTTGCTGTCGTTTTTGTGGAGAAATAGAAAAATATGTACATTCAATCCCCATGTTAAGTGGA
AGTCCCTCATAAAATTATATTAGAAATTTATTAGAAATATATACACACATATGAATATAGATAGATATACACTCTCTTT
.TTTAGCATACATAGTGCCTGTTATTTAGCAGGTACTAAAAAATAATATGTATATGTATATACATGTAGCAGAAGGCTAA
CAGGAACTTAGATACACTCAGGAATATATGATAGCATGGAAGGTTGGAACGGTGGGCCTGTGCACAATCAAGTCATGAG
GACTTAGAAAAAGACGGAACCATGGCTGGGTGCGGAGGCTCATACCTGTAATCCCAGCACTTTGGGAGGCCGAGGCCAG
TGGATTACCTGAGGTCAGGAGTTCAGGACCAGCTTGGCCAACATGGTAAAACTCTGTCTCTTAAAAAAAAAAAAAAAAA
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TGCTGTGAGCCAAGATCGCGCCATTGCAGTCCAGCCTGGGCAACAAGAGCAAAACTCAGTCTAAAAAAAAAAAAAAAAA
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GTATTTTGTTTTTGTTTTTTCTTTTTCAGACTTTGTCAACAGGAAGTAGGTGCCCATGTTGTTGATAGAAAGTTTGTAA
GTAGGACTGCACAGACTTGGTTTTGGTTCAGCATTTGGCAATAATCAGTCTTTCTGCTTCAGCCTTCAGAGAGCCCACT
TCATCATTCTTTCTGGCATCCCTGTGATCATATGATTACATATCTAATCAGTGTAACAACAACAGCTACTTAAAGAAGG
CATGCCATTAAAGATATTGTTAATATCTCTACATTGCCTTTCAAACATATGTAAGCATTCTAACTTGGAGTGAAATCTT
CTTAGGTGCTTTATATGTTGAAATTCTACCAGTCAGCTGAGGAAGAAATGTATACGGCTTATCCCCAAAATAAGATATT
CCAACAAATAATGTTTATGAGAGCTGTTAATTTATGTTAAAAAAAAAAGCTATTAATAAAATGTTTAAGTAATTAAATT
TGGGAGCTTGATTGGCAGTAGGAATATTAAGAGGATTAGCTAGATAACTAATGTAGAAAATAATATAATTGTATTGAGT
CAACAAAAGCTTATATAGATTTAATTACTATGATGATATTAGATTACTTCATAATTAGAATCTTTGTAGAATTGTTTTG
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ATCTGAAATCATATACACCAAAAGATAAGTAAAATTTGATAGAAAAAAATGAATGTATTTTAGCTGAGTTCTGACATG
CATTTTATTCACAAATCAAACTTCAGAACACAGAAGACAATTTCTTATTTTGAATTTATAGAGACTATTCTAATATTAA
ATTGATTTTTACAAGCTAGCAAATATTTTCAATAGTTGAAGCTTCGGAGTTTCTCATTTATCTAAGCTATGTAAATGCA
TGCAAAGTTTCTTCTAGAAAACAAATTACTGAAAGACATTTTCCTGATTTGTATTTGGCTGTGTTATTCCCCAGAAGGT
GAAATTATTAAACATGCCATTCGAAAGCCAGTAACTCCTTAGTACAGGTTGAACATCCCTAATCTGAAAATCTGAAATC
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ATGGGTACCAATCAAAACTTTGCTTCATGAACAAAACTATTTTAAAATATGTATAAAATTATCTTCAGGCTATTTGTTA
GGAAATAAAAATAAAACTCTAAGTCCCTCCGACTGACCCAGCGGATTCTCTCTTGGCCAAGGGAACCCCAGCAAAACCT
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TTAGGTTTTCTTCCTTAACAGCTAAACAGAAACCAGCCTTTCCAAAAGACTACTAGCTTATCTTCCCAGATACGTAACA
GTCCAGATGAGATTCATTATGTTCACTTATTTTATGTAAGATGTAGATTTACCAGGCACTAACTAAAGTTTTACAAGT
ATGTAATCATTTGTCTCACTGCTGCCCCACCACCTCCCCTGCCTTTTAAGGAAAATATATAAATACTAAACCTCCTAAG
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ACACTTCCAGTTCCAAGCATTTAGGACAAGGGATATTCAATCTGTATCAGAAACATGCGATGGTGACCACAAAAGGAAT
CTGGCAGCTGAAAATTCTGAGTCACTATCATGTCAAACTATAAAAATTTTATACAACTTAAAAAAAAAAACCTGCACTAA
TCATTAATTTATATTATTGTTAAAATTATCATACAGTAAAATTGATTTTTTTTCCTTTTGATGTACAGTTCTGAAACCA
CCTTTGCAAAAATTGCAACACTGAGAAAACTTTGACAGTGAAGAAATTTGACCTAACCAACTCCACATTGCCTTTAAC
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CTCTTTCCAAAACTAAACTGCCTTTGTAAAACTAATGAAAGGCCACCAGTTTACGAAGATAGGAGGGCCTGAATTCTGC
TAAGATATAGGCATAGTTAAGTGATTACCAGCCATTATTCCAGAGGTCACAAGATTTTCAACTTCCTCAATTACTCCTG
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TATTTGTATTAAACTCTTTCTCTACTGCAATTGCCCTATCTTGATAAATCAGCTTTATCTGAGCAGCAGGCAAGAAGAA
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CACATTCCATAAGTGTACTCTATGATGCTGATATTTATCTTTAGCCTCAGCTTATGCCAAACTATGCCTTTCCATATAT
GGCCAAÀATGTGTATTCAATATACCTCTAACTGGGAGGATCAATTGAAAGTGTTGTTTTACTAAACTCTATAACACAAA
GGAATTCAAGGATACCATAGGGATCATTCAACTGATTATAACACAAAATGGGGAGAGCCAATTTCCCTGTCCCCAGAGA
```

Fig. 6.356

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CCTCTGGGCAGCTCTAGCTCCCTGAGGGASSCAGCACAACAGTGGTCCCTTTCTCAGGAGATACCTGGGACACCAGGCA
GTTCCTCTTAGACCTTCTGGGAGAGGGGTTCCCAGCAGACGCTGGATAGGAGCAATAGCACTGAGAAGGGGTAGCAGAA
GCAGCAGTGGCCTTGACTTGTGTATGTGCACATATGCAAGGAGCCCTTGCGAGACTCCCTTCTCCCCAGGTAGTAGGGT
CTTGCTTGCCTTGCTAGTGGCAAGAACCACTAAAGTTTAGTTGTTACTGTTAACGTAACCCTATTTCTACTTACAATGT
GGTGAGGCCTTGGAAAACCCAAGTTATATAAACATTTGTCTTAGTTCATTTAGGCTGCTTTAACAAAAATACCATAAAC
TGGGTGGTTTATAAACAACGAACATTTATTTCTCACAGTTCTGGAGGCTGAGAAGTCCGAGATCATGCTGCCAGCAGAT
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TTCATTCTGATGCCCAGGTATGACAGACCTCAAATCTGTTACCGTCCCATGAACCAGGAATTATACAAAGGGTAGAGGG
GCATAGCTCTGGGTGAAGAGCTGCACTCTGTTCACTCTGGGCTCTGTGTGGTGGTGTCCCAGGTAGTGTCAGTCCAGCT
GTTAAGTCCAGTGAGGCTCAGAAAGGTGATGTCATTTGTTCAAATCTTCCAGCTAGTTTGTTGGTCAGATCATAACCAG
ACCCTATGAAGTCTAATCATACCCAGTTCTGTAGGCTTTCCTCTTCAATAGGTCCATTCTTATCTCTAACTGAGACCAC
CTGCATCCTCATAGCTTAACTATCAGTATGGAAAACAAACCATCTGTGTCACTATGACTTTACTATTTGTTCAAGAAAA
TTCTTGTCCAGGAAGATAAGACGGTGTACAATAAATAACATCAATGTTTACTTCAGAAAATTTCTGAAACCAATTTAT
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TTGCCGAATTTCCACCAGATTTCTTCCTTGAAAGGCCTGACTTTAAACCACTTGAGCTCAGACCCTTAATGTTTATAAA
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GGAAGCCTCCCCCACTGCAGTGTTAAGGTAAGTCTCTCATTGGATTTGAGTCCCGGTGTCCTTTCCATAAAGCTCTTAA
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CTTTGTGGAAATGTTTTATTACTAGTAACATTACTCTTTTTGTCTTTTATCCTATTTGTTGTGAGTCTATGAGAAGAAG
CTCACAGGGAAGAAGACAGCCGTAGACCTGGCAATTTTGTCCTCAAACTGGCTCAAAGATACAATAATATCAGCGGGTC
TCATCAAACTGGCATTCTTATAGAATAAACTTTGCTCTGGGTCACTTACTAATATCTTACAGAAAATTTATAGCAGCAG
TTGTATATTGAGGGTGTGAATAAAAACCACCAGAGAAGCTTTTTGAAAAAATACTATGAATAGATTTCTGCTTCTGGCC
AAAATGAAGAAACAGGGACCAGATTTACTCTCCAGTAAACCAGTGGAAAAAAAAACAGACACAATATTGAAAATAAAA
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```

Fig. 6.357

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Position of N ambiguity code		
30102 R	150961 Y	290063 R
30205 Y	152214 R	290164 Y
30559 Y	154374 Y	290801 R
30699 K	157074 M	292925 R
34304 R	157272 R	293201 R
34516 K	160863 Y	293611 Y
34782 R	161195 R	295755 R
35697 K	162720 Y	296143 R
35810 Y	163290 R	296739 Y
36817 Y	165441 K	297107 W
40290 K	166462 R	297460 Y
40454 M	168136 Y	297895 R
49148 S	173481 R	298027 Y
55023 Y	173519 R	298152 N
58397 Y	175259 S	298153 N
58622 R	175603 Y	298585 S
58633 S	181225 Y	298605 K
74447 R	197941 M	298799 R
75896 K	198444 Y	299792 M
82244 S	198745 R	300815 Y
88456 W	221134 R	305880 R
88499 R	222532 K	306978 M
90688 S	224195 R	309436 Y
99035 R	224801 Y	309763 Y
102977 R	226923 R	313529 K
104552 Y	227254 Y	313971 R
104862 R	227460 S	317210 S
105225 Y	228326 K	318829 Y
111252 Y	228647 Y	410826 R
111781 Y	228831 R	
112118 M	230175 K	
118914 W	230288 Y	
120628 R	232201 M	
123312 R	232338 M	
123426 S	234332 R	
125304 M	235271 R	
128015 Y	263539 K	
128393 R	270257 R	
129360 Y	270458 Y	
129361 Y	270498 R	
131865 M	271159 Y	
132562 R	274150 Y	
135112 K	274353 M	
138281 Y	275602 Y	
138806 R	277422 M	
147700 Y	278146 R	
147715 R	286615 Y	
148161 Y	289348 S	
148236 Y	289425 R	
148606 K	289868 R	
	289979 Y	

Fig. 6.358

```
<210> 2
<211> 809
<212> PRT
<213> Homo Sapien

<400> 2
Met Glu Ala Glu Gly Ser Ser Ala Pro Ala Arg Ala Gly Ser Gly Glu
1               5                   10                  15
Gly Ser Asp Ser Ala Gly Gly Ala Thr Leu Lys Ala Pro Lys His Leu
            20                  25                  30
Trp Arg His Glu Gln His His Gln Tyr Pro Leu Arg Gln Pro Gln Phe
        35                  40                  45
Arg Leu Leu His Pro His His His Leu Pro Pro Pro Pro Pro Pro Ser
    50                  55                  60
Pro Gln Pro Gln Pro Gln Cys Pro Leu Gln Pro Pro Pro Pro Pro
65                  70                  75                  80
Leu Pro Pro Pro Pro Pro Pro Gly Ala Ala Arg Gly Arg Tyr Ala
            85                  90                  95
Ser Ser Gly Ala Thr Gly Arg Val Arg His Arg Gly Tyr Ser Asp Thr
            100                 105                 110
Glu Arg Tyr Leu Tyr Cys Arg Ala Met Asp Arg Thr Ser Tyr Ala Val
            115                 120                 125
Glu Thr Gly His Arg Pro Gly Leu Lys Lys Ser Arg Met Ser Trp Pro
    130                 135                 140
Ser Ser Phe Gln Gly Leu Arg Arg Phe Asp Val Asp Asn Gly Thr Ser
145                 150                 155                 160
Ala Gly Arg Ser Pro Leu Asp Pro Met Thr Ser Pro Gly Ser Gly Leu
                165                 170                 175
Ile Leu Gln Ala Asn Phe Val His Ser Gln Arg Arg Glu Ser Phe Leu
            180                 185                 190
Tyr Arg Ser Asp Ser Asp Tyr Asp Leu Ser Pro Lys Ser Met Ser Arg
        195                 200                 205
Asn Ser Ser Ile Ala Ser Asp Ile His Gly Asp Asp Leu Ile Val Thr
    210                 215                 220
Pro Phe Ala Gln Val Leu Ala Ser Leu Arg Thr Val Arg Asn Asn Phe
225                 230                 235                 240
Ala Ala Leu Thr Asn Leu Gln Asp Arg Ala Pro Ser Lys Arg Ser Pro
                245                 250                 255
Met Cys Asn Gln Pro Ser Ile Asn Lys Ala Thr Ile Thr Glu Glu Ala
            260                 265                 270
Tyr Gln Lys Leu Ala Ser Glu Thr Leu Glu Glu Leu Asp Trp Cys Leu
        275                 280                 285
Asp Gln Leu Glu Thr Leu Gln Thr Arg His Ser Val Ser Glu Met Ala
    290                 295                 300
Ser Asn Lys Phe Lys Arg Met Leu Asn Arg Glu Leu Thr His Leu Ser
305                 310                 315                 320
Glu Met Ser Arg Ser Gly Asn Gln Val Ser Glu Phe Ile Ser Asn Thr
                325                 330                 335
Phe Leu Asp Lys Gln His Glu Val Glu Ile Pro Ser Pro Thr Gln Lys
            340                 345                 350
Glu Lys Glu Lys Lys Lys Arg Pro Met Ser Gln Ile Ser Gly Val Lys
        355                 360                 365
Lys Leu Met His Ser Ser Ser Leu Thr Asn Ser Ser Ile Pro Arg Phe
    370                 375                 380
Gly Val Lys Thr Glu Gln Glu Asp Val Leu Ala Lys Glu Leu Glu Asp
385                 390                 395                 400
Val Asn Lys Trp Gly Leu His Val Phe Arg Ile Ala Glu Leu Ser Gly
                405                 410                 415
Asn Arg Pro Leu Thr Val Ile Met His Thr Ile Phe Gln Glu Arg Asp
            420                 425                 430
Leu Leu Lys Thr Phe Lys Ile Pro Val Asp Thr Leu Ile Thr Tyr Leu
        435                 440                 445
Met Thr Leu Glu Asp His Tyr His Ala Asp Val Ala Tyr His Asn Asn
    450                 455                 460
```

Fig. 7.1
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SDOCID: <WO__02074992A2_I_>
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Ile His Ala Ala Asp Val Val Gln Ser Thr His Val Leu Leu Ser Thr
465             470             475                 480
Pro Ala Leu Glu Ala Val Phe Thr Asp Leu Glu Ile Leu Ala Ala Ile
            485             490                 495
Phe Ala Ser Ala Ile His Asp Val Asp His Pro Gly Val Ser Asn Gln
            500             505                 510
Phe Leu Ile Asn Thr Asn Ser Glu Leu Ala Leu Met Tyr Asn Asp Ser
            515             520             525

Ser Val Leu Glu Asn His His Leu Ala Val Gly Phe Lys Leu Leu Gln
        530             535             540
Glu Glu Asn Cys Asp Ile Phe Gln Asn Leu Thr Lys Lys Gln Arg Gln
545             550             555                 560
Ser Leu Arg Lys Met Val Ile Asp Ile Val Leu Ala Thr Asp Met Ser
            565             570                 575
Lys His Met Asn Leu Leu Ala Asp Leu Lys Thr Met Val Glu Thr Lys
            580             585                 590
Lys Val Thr Ser Ser Gly Val Leu Leu Leu Asp Asn Tyr Ser Asp Arg
            595             600             605
Ile Gln Val Leu Gln Asn Met Val His Cys Ala Asp Leu Ser Asn Pro
    610             615             620
Thr Lys Pro Leu Gln Leu Tyr Arg Gln Trp Thr Asp Arg Ile Met Glu
625             630             635                 640
Glu Phe Phe Arg Gln Gly Asp Arg Glu Arg Glu Arg Gly Met Glu Ile
            645             650                 655
Ser Pro Met Cys Asp Lys His Asn Ala Ser Val Glu Lys Ser Gln Val
            660             665                 670
Gly Phe Ile Asp Tyr Ile Val His Pro Leu Trp Glu Thr Trp Ala Asp
            675             680             685
Leu Val His Pro Asp Ala Gln Asp Ile Leu Asp Thr Leu Glu Asp Asn
    690             695             700
Arg Glu Trp Tyr Gln Ser Thr Ile Pro Gln Ser Pro Ser Pro Ala Pro
705             710             715                 720
Asp Asp Pro Glu Glu Gly Arg Gln Gly Gln Thr Glu Lys Phe Gln Phe
            725             730                 735
Glu Leu Thr Leu Glu Glu Asp Gly Glu Ser Asp Thr Glu Lys Asp Ser
            740             745                 750
Gly Ser Gln Val Glu Glu Asp Thr Ser Cys Ser Asp Ser Lys Thr Leu
    755             760                 765
Cys Thr Gln Asp Ser Glu Ser Thr Glu Ile Pro Leu Asp Glu Gln Val
    770             775             780
Glu Glu Glu Ala Val Gly Glu Glu Glu Ser Gln Pro Glu Ala Cys
785             790             795                 800
Val Ile Asp Asp Arg Ser Pro Asp Thr
            805
```

```
<210> 3
<211> 150
<212> PRT
<213> Homo Sapien

<400> 3
Met Asp Arg Thr Ser Tyr Ala Val Glu Thr Gly His Arg Pro Gly Leu
1               5               10              15
Lys Lys Ser Arg Met Ser Trp Pro Ser Ser Phe Gln Gly Leu Arg Arg
            20              25              30
Phe Asp Val Asp Asn Gly Thr Ser Ala Gly Arg Ser Pro Leu Asp Pro
        35              40              45
Met Thr Ser Pro Gly Ser Gly Leu Ile Leu Gln Ala Asn Phe Val His
    50              55              60
Ser Gln Arg Arg Glu Ser Phe Leu Tyr Arg Ser Asp Ser Asp Tyr Asp
65              70              75              80
Leu Ser Pro Lys Ser Met Ser Arg Asn Ser Ser Ile Ala Ser Asp Ile
            85              90              95
```

Fig. 7.2

His Gly Asp Asp Leu Ile Val Thr Pro Phe Ala Gln Val Leu Ala Ser
              100   105   110

Leu Arg Thr Val Arg Asn Asn Phe Ala Ala Leu Thr Asn Leu Gln Asp
        115   120   125
Arg Ala Pro Ser Lys Arg Ser Pro Met Cys Asn Gln Pro Ser Ile Asn
    130   135   140
Lys Ala Thr Ile Thr Val
145   150

<210> 4
<211> 745
<212> PRT
<213> Homo Sapien

<400> 4
Met Ala Gln Gln Thr Ser Pro Asp Thr Leu Thr Val Pro Glu Val Asp
  1   5   10   15
Asn Pro His Cys Pro Asn Pro Trp Leu Asn Glu Asp Leu Val Lys Ser
            20   25   30
Leu Arg Glu Asn Leu Leu Gln His Glu Lys Ser Lys Thr Ala Arg Lys
        35   40   45
Ser Val Ser Pro Lys Leu Ser Pro Val Ile Ser Pro Arg Asn Ser Pro
    50   55   60
Arg Leu Leu Arg Arg Met Leu Leu Ser Ser Asn Ile Pro Lys Gln Arg
65   70   75   80
Arg Phe Thr Val Ala His Thr Cys Phe Asp Val Asp Asn Gly Thr Ser
            85   90   95
Ala Gly Arg Ser Pro Leu Asp Pro Met Thr Ser Pro Gly Ser Gly Leu
            100   105   110
Ile Leu Gln Ala Asn Phe Val His Ser Gln Arg Arg Glu Ser Phe Leu
        115   120   125
Tyr Arg Ser Asp Ser Asp Tyr Asp Leu Ser Pro Lys Ser Met Ser Arg
    130   135   140
Asn Ser Ser Ile Ala Ser Asp Ile His Gly Asp Asp Leu Ile Val Thr
145   150   155   160
Pro Phe Ala Gln Val Leu Ala Ser Leu Arg Thr Val Arg Asn Asn Phe
                165   170   175
Ala Ala Leu Thr Asn Leu Gln Asp Arg Ala Pro Ser Lys Arg Ser Pro
            180   185   190
Met Cys Asn Gln Pro Ser Ile Asn Lys Ala Thr Ile Thr Glu Glu Ala
        195   200   205
Tyr Gln Lys Leu Ala Ser Glu Thr Leu Glu Glu Leu Asp Trp Cys Leu
    210   215   220
Asp Gln Leu Glu Thr Leu Gln Thr Arg His Ser Val Ser Glu Met Ala
225   230   235   240
Ser Asn Lys Phe Lys Arg Met Leu Asn Arg Glu Leu Thr His Leu Ser
                245   250   255
Glu Met Ser Arg Ser Gly Asn Gln Val Ser Glu Phe Ile Ser Asn Thr
            260   265   270
Phe Leu Asp Lys Gln His Glu Val Glu Ile Pro Ser Pro Thr Gln Lys
        275   280   285
Glu Lys Glu Lys Lys Lys Arg Pro Met Ser Gln Ile Ser Gly Val Lys
    290   295   300
Lys Leu Met His Ser Ser Ser Leu Thr Asn Ser Ser Ile Pro Arg Phe
305   310   315   320
Gly Val Lys Thr Glu Gln Glu Asp Val Leu Ala Lys Glu Leu Glu Asp
                325   330   335
Val Asn Lys Trp Gly Leu His Val Phe Arg Ile Ala Glu Leu Ser Gly
            340   345   350

Asn Arg Pro Leu Thr Val Ile Met His Thr Ile Phe Gln Glu Arg Asp
        355   360   365
Leu Leu Lys Thr Phe Lys Ile Pro Val Asp Thr Leu Ile Thr Tyr Leu
    370   375   380

Fig. 7.3

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```
Met Thr Leu Glu Asp His Tyr His Ala Asp Val Ala Tyr His Asn Asn
385             390             395                 400
Ile His Ala Ala Asp Val Val Gln Ser Thr His Val Leu Leu Ser Thr
            405             410                 415
Pro Ala Leu Glu Ala Val Phe Thr Asp Leu Glu Ile Leu Ala Ala Ile
            420             425                 430
Phe Ala Ser Ala Ile His Asp Val Asp His Pro Gly Val Ser Asn Gln
            435             440                 445
Phe Leu Ile Asn Thr Asn Ser Glu Leu Ala Leu Met Tyr Asn Asp Ser
    450             455             460
Ser Val Leu Glu Asn His His Leu Ala Val Gly Phe Lys Leu Leu Gln
465             470             475                 480
Glu Glu Asn Cys Asp Ile Phe Gln Asn Leu Thr Lys Lys Gln Arg Gln
            485             490                 495
Ser Leu Arg Lys Met Val Ile Asp Ile Val Leu Ala Thr Asp Met Ser
            500             505                 510
Lys His Met Asn Leu Leu Ala Asp Leu Lys Thr Met Val Glu Thr Lys
            515             520                 525
Lys Val Thr Ser Ser Gly Val Leu Leu Leu Asp Asn Tyr Ser Asp Arg
    530             535             540
Ile Gln Val Leu Gln Asn Met Val His Cys Ala Asp Leu Ser Asn Pro
545             550             555                 560
Thr Lys Pro Leu Gln Leu Tyr Arg Gln Trp Thr Asp Arg Ile Met Glu
            565             570                 575
Glu Phe Phe Arg Gln Gly Asp Arg Glu Arg Glu Arg Gly Met Glu Ile
            580             585                 590
Ser Pro Met Cys Asp Lys His Asn Ala Ser Val Glu Lys Ser Gln Val
            595             600                 605
Gly Phe Ile Asp Tyr Ile Val His Pro Leu Trp Glu Thr Trp Ala Asp
    610             615             620
Leu Val His Pro Asp Ala Gln Asp Ile Leu Asp Thr Leu Glu Asp Asn
625             630             635                 640
Arg Glu Trp Tyr Gln Ser Thr Ile Pro Gln Ser Pro Ser Pro Ala Pro
            645             650                 655
Asp Asp Pro Glu Glu Gly Arg Gln Gly Gln Thr Glu Lys Phe Gln Phe
            660             665                 670
Glu Leu Thr Leu Glu Glu Asp Gly Glu Ser Asp Thr Glu Lys Asp Ser
    675             680             685
Gly Ser Gln Val Glu Glu Asp Thr Ser Cys Ser Asp Ser Lys Thr Leu
    690             695             700
Cys Thr Gln Asp Ser Glu Ser Thr Glu Ile Pro Leu Asp Glu Gln Val
705             710             715                 720
Glu Glu Glu Ala Val Gly Glu Glu Glu Glu Ser Gln Pro Glu Ala Cys
            725             730                 735
Val Ile Asp Asp Arg Ser Pro Asp Thr
            740             745
```

<210> 5
<211> 215
<212> PRT
<213> Homo Sapien

<400> 5

```
Met Ala Gln Gln Thr Ser Pro Asp Thr Leu Thr Val Pro Glu Val Asp
1               5               10              15
Asn Pro His Cys Pro Asn Pro Trp Leu Asn Glu Asp Leu Val Lys Ser
            20              25              30
Leu Arg Glu Asn Leu Leu Gln His Glu Lys Ser Lys Thr Ala Arg Lys
    35              40              45
Ser Val Ser Pro Lys Leu Ser Pro Val Ile Ser Pro Arg Asn Ser Pro
    50              55              60
Arg Leu Leu Arg Arg Met Leu Leu Ser Ser Asn Ile Pro Lys Gln Arg
65              70              75                  80
```

Fig. 7.4

```
Arg Phe Thr Val Ala His Thr Cys Phe Asp Val Asp Asn Gly Thr Ser
            85                  90                  95
Ala Gly Arg Ser Pro Leu Asp Pro Met Thr Ser Pro Gly Ser Gly Leu
            100                 105                 110
Ile Leu Gln Ala Asn Phe Val His Ser Gln Arg Arg Glu Ser Phe Leu
            115                 120                 125
Tyr Arg Ser Asp Ser Asp Tyr Asp Leu Ser Pro Lys Ser Met Ser Arg
    130                 135                 140
Asn Ser Ser Ile Ala Ser Asp Ile His Gly Asp Asp Leu Ile Val Thr
145             150                 155                 160
Pro Phe Ala Gln Val Leu Ala Ser Leu Arg Thr Val Arg Asn Asn Phe
            165                 170                 175
Ala Ala Leu Thr Asn Leu Gln Asp Arg Ala Pro Ser Lys Arg Ser Pro
            180                 185                 190
Met Cys Asn Gln Pro Ser Ile Asn Lys Ala Thr Ile Thr Gly Leu Tyr
        195                 200                 205
Asn Gly Ile Ile Ala Phe Leu
    210                 215
```

<210> 6
<211> 673
<212> PRT
<213> Homo Sapien

<400> 6

```
Met Met His Val Asn Asn Phe Pro Phe Arg Arg His Ser Trp Ile Cys
1               5                   10                  15
Phe Asp Val Asp Asn Gly Thr Ser Ala Gly Arg Ser Pro Leu Asp Pro
            20                  25                  30
Met Thr Ser Pro Gly Ser Gly Leu Ile Leu Gln Ala Asn Phe Val His
        35                  40                  45
Ser Gln Arg Arg Glu Ser Phe Leu Tyr Arg Ser Asp Ser Asp Tyr Asp
    50                  55                  60
Leu Ser Pro Lys Ser Met Ser Arg Asn Ser Ser Ile Ala Ser Asp Ile
65              70                  75                  80
His Gly Asp Asp Leu Ile Val Thr Pro Phe Ala Gln Val Leu Ala Ser
            85                  90                  95
Leu Arg Thr Val Arg Asn Asn Phe Ala Ala Leu Thr Asn Leu Gln Asp
            100                 105                 110
Arg Ala Pro Ser Lys Arg Ser Pro Met Cys Asn Gln Pro Ser Ile Asn
    115                 120                 125
Lys Ala Thr Ile Thr Glu Glu Ala Tyr Gln Lys Leu Ala Ser Glu Thr
    130                 135                 140
Leu Glu Glu Leu Asp Trp Cys Leu Asp Gln Leu Glu Thr Leu Gln Thr
145             150                 155                 160
Arg His Ser Val Ser Glu Met Ala Ser Asn Lys Phe Lys Arg Met Leu
                165                 170                 175

Asn Arg Glu Leu Thr His Leu Ser Glu Met Ser Arg Ser Gly Asn Gln
            180                 185                 190
Val Ser Glu Phe Ile Ser Asn Thr Phe Leu Asp Lys Gln His Glu Val
        195                 200                 205
Glu Ile Pro Ser Pro Thr Gln Lys Glu Lys Glu Lys Lys Lys Arg Pro
    210                 215                 220
Met Ser Gln Ile Ser Gly Val Lys Lys Leu Met His Ser Ser Ser Leu
225             230                 235                 240
Thr Asn Ser Ser Ile Pro Arg Phe Gly Val Lys Thr Glu Gln Glu Asp
            245                 250                 255
Val Leu Ala Lys Glu Leu Glu Asp Val Asn Lys Trp Gly Leu His Val
            260                 265                 270
Phe Arg Ile Ala Glu Leu Ser Gly Asn Arg Pro Leu Thr Val Ile Met
    275                 280                 285
His Thr Ile Phe Gln Glu Arg Asp Leu Leu Lys Thr Phe Lys Ile Pro
    290                 295                 300
```

Fig. 7.5

```
Val Asp Thr Leu Ile Thr Tyr Leu Met Thr Leu Glu Asp His Tyr His
305                 310                 315                 320
Ala Asp Val Ala Tyr His Asn Asn Ile His Ala Ala Asp Val Val Gln
                325             330             335
Ser Thr His Val Leu Leu Ser Thr Pro Ala Leu Glu Ala Val Phe Thr
        340             345             350
Asp Leu Glu Ile Leu Ala Ala Ile Phe Ala Ser Ala Ile His Asp Val
        355             360             365
Asp His Pro Gly Val Ser Asn Gln Phe Leu Ile Asn Thr Asn Ser Glu
    370             375             380
Leu Ala Leu Met Tyr Asn Asp Ser Ser Val Leu Glu Asn His His Leu
385             390             395             400
Ala Val Gly Phe Lys Leu Leu Gln Glu Glu Asn Cys Asp Ile Phe Gln
                405             410             415
Asn Leu Thr Lys Lys Gln Arg Gln Ser Leu Arg Lys Met Val Ile Asp
            420             425             430
Ile Val Leu Ala Thr Asp Met Ser Lys His Met Asn Leu Leu Ala Asp
            435             440             445
Leu Lys Thr Met Val Glu Thr Lys Lys Val Thr Ser Ser Gly Val Leu
    450             455             460
Leu Leu Asp Asn Tyr Ser Asp Arg Ile Gln Val Leu Gln Asn Met Val
465             470             475             480
His Cys Ala Asp Leu Ser Asn Pro Thr Lys Pro Leu Gln Leu Tyr Arg
                485             490             495
Gln Trp Thr Asp Arg Ile Met Glu Glu Phe Phe Arg Gln Gly Asp Arg
            500             505             510
Glu Arg Glu Arg Gly Met Glu Ile Ser Pro Met Cys Asp Lys His Asn
    515             520             525
Ala Ser Val Glu Lys Ser Gln Val Gly Phe Ile Asp Tyr Ile Val His
    530             535             540
Pro Leu Trp Glu Thr Trp Ala Asp Leu Val His Pro Asp Ala Gln Asp
545             550             555             560
Ile Leu Asp Thr Leu Glu Asp Asn Arg Glu Trp Tyr Gln Ser Thr Ile
            565             570             575
Pro Gln Ser Pro Ser Pro Ala Pro Asp Asp Pro Glu Glu Gly Arg Gln
        580             585             590
Gly Gln Thr Glu Lys Phe Gln Phe Glu Leu Thr Leu Glu Glu Asp Gly
    595             600             605
Glu Ser Asp Thr Glu Lys Asp Ser Gly Ser Gln Val Glu Glu Asp Thr
    610             615             620
Ser Cys Ser Asp Ser Lys Thr Leu Cys Thr Gln Asp Ser Glu Ser Thr
625             630             635             640

Glu Ile Pro Leu Asp Glu Gln Val Glu Glu Glu Ala Val Gly Glu Glu
            645             650             655
Glu Glu Ser Gln Pro Glu Ala Cys Val Ile Asp Asp Arg Ser Pro Asp
        660             665             670
Thr
```

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<210> 7
<211> 15
<212> PRT
<213> Homo Sapien

<400> 7
Met Met His Val Asn Asn Phe Pro Phe Arg Arg His Ser Trp Ile
1                 5                 10                  15


<210> 8
<211> 687
<212> PRT
<213> Homo Sapien

<400> 8
```

Fig. 7.6

```
Met Ala Phe Val Trp Asp Pro Leu Gly Ala Thr Val Pro Gly Pro Ser
1               5                   10                  15
Thr Arg Ala Lys Ser Arg Leu Arg Phe Ser Lys Ser Tyr Ser Phe Asp
            20                  25                  30
Val Asp Asn Gly Thr Ser Ala Gly Arg Ser Pro Leu Asp Pro Met Thr
            35                  40                  45
Ser Pro Gly Ser Gly Leu Ile Leu Gln Ala Asn Phe Val His Ser Gln
    50                  55                  60
Arg Arg Glu Ser Phe Leu Tyr Arg Ser Asp Ser Asp Tyr Asp Leu Ser
65                  70                  75                  80
Pro Lys Ser Met Ser Arg Asn Ser Ser Ile Ala Ser Asp Ile His Gly
            85                  90                  95
Asp Asp Leu Ile Val Thr Pro Phe Ala Gln Val Leu Ala Ser Leu Arg
            100                 105                 110
Thr Val Arg Asn Asn Phe Ala Ala Leu Thr Asn Leu Gln Asp Arg Ala
        115                 120                 125
Pro Ser Lys Arg Ser Pro Met Cys Asn Gln Pro Ser Ile Asn Lys Ala
    130                 135                 140
Thr Ile Thr Glu Glu Ala Tyr Gln Lys Leu Ala Ser Glu Thr Leu Glu
145                 150                 155                 160
Glu Leu Asp Trp Cys Leu Asp Gln Leu Glu Thr Leu Gln Thr Arg His
                165                 170                 175
Ser Val Ser Glu Met Ala Ser Asn Lys Phe Lys Arg Met Leu Asn Arg
            180                 185                 190
Glu Leu Thr His Leu Ser Glu Met Ser Arg Ser Gly Asn Gln Val Ser
        195                 200                 205
Glu Phe Ile Ser Asn Thr Phe Leu Asp Lys Gln His Glu Val Glu Ile
    210                 215                 220
Pro Ser Pro Thr Gln Lys Glu Lys Glu Lys Lys Lys Arg Pro Met Ser
225                 230                 235                 240
Gln Ile Ser Gly Val Lys Lys Leu Met His Ser Ser Ser Leu Thr Asn
            245                 250                 255
Ser Ser Ile Pro Arg Phe Gly Val Lys Thr Glu Gln Glu Asp Val Leu
            260                 265                 270

Ala Lys Glu Leu Glu Asp Val Asn Lys Trp Gly Leu His Val Phe Arg
        275                 280                 285
Ile Ala Glu Leu Ser Gly Asn Arg Pro Leu Thr Val Ile Met His Thr
    290                 295                 300
Ile Phe Gln Glu Arg Asp Leu Leu Lys Thr Phe Lys Ile Pro Val Asp
305                 310                 315                 320
Thr Leu Ile Thr Tyr Leu Met Thr Leu Glu Asp His Tyr His Ala Asp
                325                 330                 335
Val Ala Tyr His Asn Asn Ile His Ala Ala Asp Val Val Gln Ser Thr
            340                 345                 350
His Val Leu Leu Ser Thr Pro Ala Leu Glu Ala Val Phe Thr Asp Leu
        355                 360                 365
Glu Ile Leu Ala Ala Ile Phe Ala Ser Ala Ile His Asp Val Asp His
    370                 375                 380
Pro Gly Val Ser Asn Gln Phe Leu Ile Asn Thr Asn Ser Glu Leu Ala
385                 390                 395                 400
Leu Met Tyr Asn Asp Ser Ser Val Leu Glu Asn His His Leu Ala Val
                405                 410                 415
Gly Phe Lys Leu Leu Gln Glu Glu Asn Cys Asp Ile Phe Gln Asn Leu
            420                 425                 430
Thr Lys Lys Gln Arg Gln Ser Leu Arg Lys Met Val Ile Asp Ile Val
        435                 440                 445
Leu Ala Thr Asp Met Ser Lys His Met Asn Leu Leu Ala Asp Leu Lys
    450                 455                 460
Thr Met Val Glu Thr Lys Lys Val Thr Ser Ser Gly Val Leu Leu Leu
465                 470                 475                 480
Asp Asn Tyr Ser Asp Arg Ile Gln Val Leu Gln Asn Met Val His Cys
            485                 490                 495
Ala Asp Leu Ser Asn Pro Thr Lys Pro Leu Gln Leu Tyr Arg Gln Trp
            500                 505                 510
Thr Asp Arg Ile Met Glu Glu Phe Phe Arg Gln Gly Asp Arg Glu Arg
        515                 520                 525
```

Fig. 7.7

```
Glu Arg Gly Met Glu Ile Ser Pro Met Cys Asp Lys His Asn Ala Ser
    530                 535                 540
Val Glu Lys Ser Gln Val Gly Phe Ile Asp Tyr Ile Val His Pro Leu
545                 550                 555                 560
Trp Glu Thr Trp Ala Asp Leu Val His Pro Asp Ala Gln Asp Ile Leu
            565                 570                 575
Asp Thr Leu Glu Asp Asn Arg Glu Trp Tyr Gln Ser Thr Ile Pro Gln
            580                 585                 590
Ser Pro Ser Pro Ala Pro Asp Asp Pro Glu Glu Gly Arg Gln Gly Gln
        595                 600                 605
Thr Glu Lys Phe Gln Phe Glu Leu Thr Leu Glu Glu Asp Gly Glu Ser
    610                 615                 620
Asp Thr Glu Lys Asp Ser Gly Ser Gln Val Glu Glu Asp Thr Ser Cys
625                 630                 635                 640
Ser Asp Ser Lys Thr Leu Cys Thr Gln Asp Ser Glu Ser Thr Glu Ile
            645                 650                 655
Pro Leu Asp Glu Gln Val Glu Glu Glu Ala Val Gly Glu Glu Glu Glu
            660                 665                 670
Ser Gln Pro Glu Ala Cys Val Ile Asp Asp Arg Ser Pro Asp Thr
        675                 680                 685
```

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<210> 9
<211> 585
<212> PRT
<213> Homo Sapien

<400> 9
Met Lys Glu Gln Pro Ser Cys Ala Gly Thr Gly His Pro Ser Met Ala
1               5               10                  15
Gly Tyr Gly Arg Met Ala Pro Phe Glu Leu Ala Ser Gly Pro Val Lys
            20                  25                  30
Arg Leu Arg Thr Glu Ser Pro Phe Pro Cys Leu Phe Ala Glu Glu Ala
        35                  40                  45
Tyr Gln Lys Leu Ala Ser Glu Thr Leu Glu Glu Leu Asp Trp Cys Leu
    50                  55                  60
Asp Gln Leu Glu Thr Leu Gln Thr Arg His Ser Val Ser Glu Met Ala
65                  70                  75                  80
Ser Asn Lys Phe Lys Arg Met Leu Asn Arg Glu Leu Thr His Leu Ser
            85                  90                  95
Glu Met Ser Arg Ser Gly Asn Gln Val Ser Glu Phe Ile Ser Asn Thr
            100                 105                 110
Phe Leu Asp Lys Gln His Glu Val Glu Ile Pro Ser Pro Thr Gln Lys
        115                 120                 125
Glu Lys Glu Lys Lys Lys Arg Pro Met Ser Gln Ile Ser Gly Val Lys
    130                 135                 140
Lys Leu Met His Ser Ser Ser Leu Thr Asn Ser Ser Ile Pro Arg Phe
145                 150                 155                 160
Gly Val Lys Thr Glu Gln Glu Asp Val Leu Ala Lys Glu Leu Glu Asp
            165                 170                 175
Val Asn Lys Trp Gly Leu His Val Phe Arg Ile Ala Glu Leu Ser Gly
            180                 185                 190
Asn Arg Pro Leu Thr Val Ile Met His Thr Ile Phe Gln Glu Arg Asp
        195                 200                 205
Leu Leu Lys Thr Phe Lys Ile Pro Val Asp Thr Leu Ile Thr Tyr Leu
    210                 215                 220
Met Thr Leu Glu Asp His Tyr His Ala Asp Val Ala Tyr His Asn Asn
225                 230                 235                 240
Ile His Ala Ala Asp Val Val Gln Ser Thr His Val Leu Leu Ser Thr
            245                 250                 255
Pro Ala Leu Glu Ala Val Phe Thr Asp Leu Glu Ile Leu Ala Ala Ile
            260                 265                 270
Phe Ala Ser Ala Ile His Asp Val Asp His Pro Gly Val Ser Asn Gln
        275                 280                 285
Phe Leu Ile Asn Thr Asn Ser Glu Leu Ala Leu Met Tyr Asn Asp Ser
    290                 295                 300
```

Fig. 7.8

```
Ser Val Leu Glu Asn His His Leu Ala Val Gly Phe Lys Leu Leu Gln
305                 310                 315                 320
Glu Glu Asn Cys Asp Ile Phe Gln Asn Leu Thr Lys Lys Gln Arg Gln
                325                 330                 335
Ser Leu Arg Lys Met Val Ile Asp Ile Val Leu Ala Thr Asp Met Ser
            340                 345                 350
Lys His Met Asn Leu Leu Ala Asp Leu Lys Thr Met Val Glu Thr Lys
        355                 360                 365
Lys Val Thr Ser Ser Gly Val Leu Leu Leu Asp Asn Tyr Ser Asp Arg
    370                 375                 380
Ile Gln Val Leu Gln Asn Met Val His Cys Ala Asp Leu Ser Asn Pro
385                 390                 395                 400
Thr Lys Pro Leu Gln Leu Tyr Arg Gln Trp Thr Asp Arg Ile Met Glu
                405                 410                 415
Glu Phe Phe Arg Gln Gly Asp Arg Glu Arg Glu Arg Gly Met Glu Ile
            420                 425                 430
Ser Pro Met Cys Asp Lys His Asn Ala Ser Val Glu Lys Ser Gln Val
        435                 440                 445
Gly Phe Ile Asp Tyr Ile Val His Pro Leu Trp Glu Thr Trp Ala Asp
    450                 455                 460
Leu Val His Pro Asp Ala Gln Asp Ile Leu Asp Thr Leu Glu Asp Asn
465                 470                 475                 480
Arg Glu Trp Tyr Gln Ser Thr Ile Pro Gln Ser Pro Ser Pro Ala Pro
                485                 490                 495
Asp Asp Pro Glu Glu Gly Arg Gln Gly Gln Thr Glu Lys Phe Gln Phe
            500                 505                 510
Glu Leu Thr Leu Glu Glu Asp Gly Glu Ser Asp Thr Glu Lys Asp Ser
        515                 520                 525
Gly Ser Gln Val Glu Glu Asp Thr Ser Cys Ser Asp Ser Lys Thr Leu
    530                 535                 540
Cys Thr Gln Asp Ser Glu Ser Thr Glu Ile Pro Leu Asp Glu Gln Val
545                 550                 555                 560
Glu Glu Glu Ala Val Gly Glu Glu Glu Glu Ser Gln Pro Glu Ala Cys
                565                 570                 575
Val Ile Asp Asp Arg Ser Pro Asp Thr
            580                 585
```

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<210> 10
<211> 507
<212> PRT
<213> Homo Sapien

<400> 10
Met Ala Ser Asn Lys Phe Lys Arg Met Leu Asn Arg Glu Leu Thr His
1               5                   10                  15
Leu Ser Glu Met Ser Arg Ser Gly Asn Gln Val Ser Glu Phe Ile Ser
            20                  25                  30
Asn Thr Phe Leu Asp Lys Gln His Glu Val Glu Ile Pro Ser Pro Thr
        35                  40                  45
Gln Lys Glu Lys Glu Lys Lys Lys Arg Pro Met Ser Gln Ile Ser Gly
    50                  55                  60
Val Lys Lys Leu Met His Ser Ser Ser Leu Thr Asn Ser Ser Ile Pro
65              70                  75                  80
Arg Phe Gly Val Lys Thr Glu Gln Glu Asp Val Leu Ala Lys Glu Leu
                85                  90                  95
Glu Asp Val Asn Lys Trp Gly Leu His Val Phe Arg Ile Ala Glu Leu
            100                 105                 110
Ser Gly Asn Arg Pro Leu Thr Val Ile Met His Thr Ile Phe Gln Glu
        115                 120                 125
Arg Asp Leu Leu Lys Thr Phe Lys Ile Pro Val Asp Thr Leu Ile Thr
    130                 135                 140
Tyr Leu Met Thr Leu Glu Asp His Tyr His Ala Asp Val Ala Tyr His
145                 150                 155                 160
Asn Asn Ile His Ala Ala Asp Val Val Gln Ser Thr His Val Leu Leu
                165                 170                 175
```

Fig. 7.9

```
Ser Thr Pro Ala Leu Glu Ala Val Phe Thr Asp Leu Glu Ile Leu Ala
            180                     185                 190
Ala Ile Phe Ala Ser Ala Ile His Asp Val Asp His Pro Gly Val Ser
        195                 200                 205
Asn Gln Phe Leu Ile Asn Thr Asn Ser Glu Leu Ala Leu Met Tyr Asn
        210                 215                 220
Asp Ser Ser Val Leu Glu Asn His His Leu Ala Val Gly Phe Lys Leu
225                 230                 235                 240
Leu Gln Glu Glu Asn Cys Asp Ile Phe Gln Asn Leu Thr Lys Lys Gln
                245                 250                 255
Arg Gln Ser Leu Arg Lys Met Val Ile Asp Ile Val Leu Ala Thr Asp
            260                 265                 270

Met Ser Lys His Met Asn Leu Leu Ala Asp Leu Lys Thr Met Val Glu
            275                 280                 285
Thr Lys Lys Val Thr Ser Ser Gly Val Leu Leu Leu Asp Asn Tyr Ser
        290                 295                 300
Asp Arg Ile Gln Val Leu Gln Asn Met Val His Cys Ala Asp Leu Ser
305                 310                 315                 320
Asn Pro Thr Lys Pro Leu Gln Leu Tyr Arg Gln Trp Thr Asp Arg Ile
            325                 330                 335
Met Glu Glu Phe Phe Arg Gln Gly Asp Arg Glu Arg Glu Arg Gly Met
            340                 345                 350
Glu Ile Ser Pro Met Cys Asp Lys His Asn Ala Ser Val Glu Lys Ser
            355                 360                 365
Gln Val Gly Phe Ile Asp Tyr Ile Val His Pro Leu Trp Glu Thr Trp
        370                 375                 380
Ala Asp Leu Val His Pro Asp Ala Gln Asp Ile Leu Asp Thr Leu Glu
385                 390                 395                 400
Asp Asn Arg Glu Trp Tyr Gln Ser Thr Ile Pro Gln Ser Pro Ser Pro
            405                 410                 415
Ala Pro Asp Asp Pro Glu Glu Gly Arg Gln Gly Gln Thr Glu Lys Phe
            420                 425                 430
Gln Phe Glu Leu Thr Leu Glu Glu Asp Gly Glu Ser Asp Thr Glu Lys
        435                 440                 445
Asp Ser Gly Ser Gln Val Glu Glu Asp Thr Ser Cys Ser Asp Ser Lys
450                 455                 460
Thr Leu Cys Thr Gln Asp Ser Glu Ser Thr Glu Ile Pro Leu Asp Glu
465                 470                 475                 480
Gln Val Glu Glu Glu Ala Val Gly Glu Glu Glu Glu Ser Gln Pro Glu
            485                 490                 495
Ala Cys Val Ile Asp Asp Arg Ser Pro Asp Thr
            500                 505
```

Fig. 7.10

mRNA/cDNA variants	Isoform	142207 142328 4D7-1	444645 444775 4D7-2	641649 641878 4D7-3	736254 737226 4D4	861791 862202 4D5	1044051 1044190 4D3	1273404 1273709 4D6	1354347 1355128 4D8	1414511 1414702 LF1	1436943 1436979 LF2	1445217 1445290 LF3
UO2882	4D4									*	*	*
L20969	4D5				*					*	*	*
AF012073	4D3									*	*	*
L20970	4D2					*				*	*	*
AF012074	4D3						*			*	*	*
U50159	4D2						*					
U50158	4D1									*	*	*
U50157	4DN3											
AJ250854	4D4				*	*				*	*	*
NM_006203	4DN1									*	*	*
AJ250852	4DN2				*	*						
AJ250855	4DN3							*				
BC008390									*			
novel cDNA identified by deCODE												
RT-PCR	4D6	*	*	*						*	*	*
CAP-RACE	4D7									*	*	*
CAP-RACE	4D8								*	*	*	*

Fig. 8A

Fig. 8B

(54) Title: PHOSPHODIESTERASE 4D GENES RELATED TO HUMAN STROKE

(57) Abstract: A role of the human PDE4D gene in stroke is disclosed. New exons, referred to as 4D7-1, 4D7-2, 4D7-3, 4D6 and 4D have been identified. Moreover, three splice variants have been identified. Methods for diagnosis, predictions of clinical course and treatment for stroke using polymorphisms in the PDE4D gene are also disclosed.

For two-letter codes and other abbreviations, refer to the "Guidance Notes on Codes and Abbreviations" appearing at the beginning of each regular issue of the PCT Gazette.

INTERNATIONAL SEARCH REPORT

A. CLASSIFICATION OF SUBJECT MATTER
IPC 7   C12N9/16   C12Q1/68   C12N15/52   A61K38/46

According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)
IPC 7   C12N   C12Q   A61K

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practical, search terms used)

EPO-Internal

C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	GRAEME B BOLGER ET AL: "Characterization of five different proteins produced by alternatively spliced mRNAs from the human cAMP-specific phosphodiesterase PDE4D gene" BIOCHEMICAL JOURNAL, PORTLAND PRESS, LONDON, GB, vol. 328, 1997, pages 539-548, XP002150449 ISSN: 0264-6021 the whole document	1-18,22, 26,30, 32,34, 35,39, 40,42, 43,47-50
A		19-21,59

	-/--	

| X | Further documents are listed in the continuation of box C. |

| X | Patent family members are listed in annex. |

* Special categories of cited documents :

"A" document defining the general state of the art which is not considered to be of particular relevance

"E" earlier document but published on or after the international filing date

"L" document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)

"O" document referring to an oral disclosure, use, exhibition or other means

"P" document published prior to the international filing date but later than the priority date claimed

"T" later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention

"X" document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone

"Y" document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art.

"&" document member of the same patent family

Date of the actual completion of the international search	Date of mailing of the international search report
23 October 2002	05. 03. 2003

Name and mailing address of the ISA	Authorized officer
European Patent Office, P.B. 5818 Patentlaan 2 NL - 2280 HV Rijswijk Tel. (+31-70) 340-2040, Tx. 31 651 epo nl, Fax: (+31-70) 340-3016	Patrick Andersson

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Form PCT/ISA/210 (second sheet) (July 1992)

page 1 of 2

C.(Continuation) DOCUMENTS CONSIDERED TO BE RELEVANT		
Category°	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	XAVIER MIRÓ ET AL: "Phosphodiesterases 4D and 7A splice variants in the response of HUVEC cells to TNF-alphal." BIOCHEMICAL AND BIOPHYSICAL RESEARCH COMMUNICATIONS, vol. 274, 2000, pages 415-421, XP002902795 ACADEMIC PRESS ISSN: 0006-291X the whole document	1-18,22, 26,30, 32,34, 35,39, 40,42, 43,47-58
A	---	19,20,59
X	WO 01 00851 A (MEMORY PHARMACEUTICA! CORP) 4 January 2001 (2001-01-04) the whole document	1-18,22, 26,30, 32,34, 35,39, 40,42, 43,47-58
A	---	19-21,59
X	WO 00 23091 A (SCUDDER KURT MARSHALL ;BIOIMAGE A S (DK); THASTRUP OLE (DK); BJOER) 27 April 2000 (2000-04-27) the whole document	1-18,22, 26,30, 32,34, 35,39, 40,42, 43,47-58
A	---	19-21,59
X	WO 00 40714 A (ARROW AMY ;OLIGOS ETC INC (US); THOMPSON TERRY (US); DALE RODERIC) 13 July 2000 (2000-07-13) the whole document	30, 48-51, 54-58
A	---	59
A	WO 00 77226 A (KAPELLER LIBERMANN ROSANA ;WHITE DAVID (US); ROBISON KEITH E (US);) 21 December 2000 (2000-12-21) the whole document -----	1-22,26, 30,32, 34,35, 39,40, 42,43, 47-59

2

INTERNATIONAL SEARCH REPORT

Box I	**Observations where certain claims were found unsearchable (Continuation of item 1 of first sheet)**

This International Search Report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

1. [X] Claims Nos.:   **19, 44-46**
because they relate to subject matter not required to be searched by this Authority, namely:

    see FURTHER INFORMATION sheet PCT/ISA/210

2. [X] Claims Nos.:
because they relate to parts of the International Application that do not comply with the prescribed requirements to such an extent that no meaningful International Search can be carried out, specifically:

    see FURTHER INFORMATION sheet PCT/ISA/210

3. [ ] Claims Nos.:
because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).

Box II	**Observations where unity of invention is lacking (Continuation of item 2 of first sheet)**

This International Searching Authority found multiple inventions in this international application, as follows:

    see additional sheet

1. [ ] As all required additional search fees were timely paid by the applicant, this International Search Report covers all searchable claims.

2. [ ] As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.

3. [ ] As only some of the required additional search fees were timely paid by the applicant, this International Search Report covers only those claims for which fees were paid, specifically claims Nos.:

4. [X] No required additional search fees were timely paid by the applicant. Consequently, this International Search Report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:

    1-59 (partially)

Remark on Protest
    [ ] The additional search fees were accompanied by the applicant's protest.
    [ ] No protest accompanied the payment of additional search fees.

Form PCT/ISA/210 (continuation of first sheet (1)) (July 1998)

FURTHER INFORMATION CONTINUED FROM   PCT/ISA/ 210

This International Searching Authority found multiple (groups of) inventions in this international application, as follows:

1. Claims: 1-59 (partially)

    Each sequence of SEQ ID 1-10 and 12 represent one invention.

2. Claims: 1-59 (partially)

    Each sequence of SEQ ID 1-10 and 12 represent one invention.

3. Claims: 1-59 (partially)

    Each sequence of SEQ ID 1-10 and 12 represent one invention.

FURTHER INFORMATION CONTINUED FROM   PCT/ISA/ 210

Continuation of Box I.1

Claims Nos.: 19, 44-46

Claims 19, 44-46 relate to methods of treatment of the human or animal body by surgery or by therapy / diagnostic methods practised on the human or animal body / Rule 39.1(iv). Nevertheless, a search has been executed for these claims. The search has been based on the alleged effects of the compounds/ compositions.

Continuation of Box I.2

Claims Nos.: 23-25,27-29,31, 33, 36-38, 41, 44-46 and parts of 40 and 42.

Claims 23-25, 27-29, 31, 33, 36-38, 41, 44-46 and parts of claims 40 and 42 relate to agents interacting with a polypeptide encoded by a phosphodiesterase 4D gene or the expression of this gene.
These claims could include known compounds e.g. known phosphodiesterase inhibitors. Moreover, the description does not give any example of such substance. Identification of agents with the claimed methods does not give the identified agents PER SE any unique propertiess and thus, the description lacks disclosure and the claim lacks support within the meaning of PCT Articles 5 and 6.
A meaningful search of claims 23-25, 27-29, 31, 33, 36-38, 41, 44-46 and parts of claims 40 and 42 is impossible and consequently, the claims have not been searched.

The following parts of claims 40 and 42 have been searched: A phosphodiesterase 4D gene PER SE; fragments, variants or derivates is considered to be unclear, e.g. fragment could in its extreme be one single nucleotide.

The applicant's attention is drawn to the fact that claims, or parts of claims, relating to inventions in respect of which no international search report has been established need not be the subject of an international preliminary examination (Rule 66.1(e) PCT). The applicant is advised that the EPO policy when acting as an International Preliminary Examining Authority is normally not to carry out a preliminary examination on matter which has not been searched. This is the case irrespective of whether or not the claims are amended following receipt of the search report or during any Chapter II procedure.

INTERNATIONAL SEARCH REPORT

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